



# **PRECISION MEDICINE & GENETICS: IMPLICATIONS FOR THE ADVANCED PRACTICE PROVIDER**

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# Objectives

- Understand what precision medicine is and how it can be applied to the transplant patient.
- Identify historic and current challenges to managing immunosuppression.
- Evaluate certain groups of patients that pose challenges in being immunosuppressed.
- Discuss specific testing used in precision medicine and how to utilize the test in the transplant patient.
- Understand pharmacogenomic testing and how this can be applied to immunosuppression.
- Identify ways in which the Advanced Care Provider can use precision medicine in the care of the transplant recipient.

# What is Precision Medicine?

- Human Genome Project (1990-2003) by the U.S. Department of Energy and National Institute of Health- human blueprint. Accounted for over 90% of the human genome (what makes up a human).
- Genome-complete set of DNA. Humans have one genome which contains 23 pairs of chromosomes and 3 billion base pairs of DNA.
- The Human Genome Project is the foundation for modern precision medicine.
- Movement to: Predictive, Preventive, Personalized, Participatory Care.
- Pioneered in Oncology.

# Advancements in Transplant

"Where we have been..?"

- Between 1980's-2000's we have made progress in immunosuppression
  - Cyclosporine  Tacrolimus
  - Azathioprine  Mycophenolate
  - Steroids  Steroid Sparing
- Change in how we treat infections:
  - Ganciclovir/Foscarnet  Valganciclovir/Letermovir
- Sirolimus (1999), Everolimus (2010)
- Atgam (equine-1974)  Thymoglobulin (rabbit-1998)
- Polyclonal Antibodies (Atgam/Thymoglobulin)  Monoclonal Antibodies (Rituximab/Basiliximab)
- Reduced acute rejection rates and improved survival within the first year.

# Advancements in Transplant

"...and where we are today?"

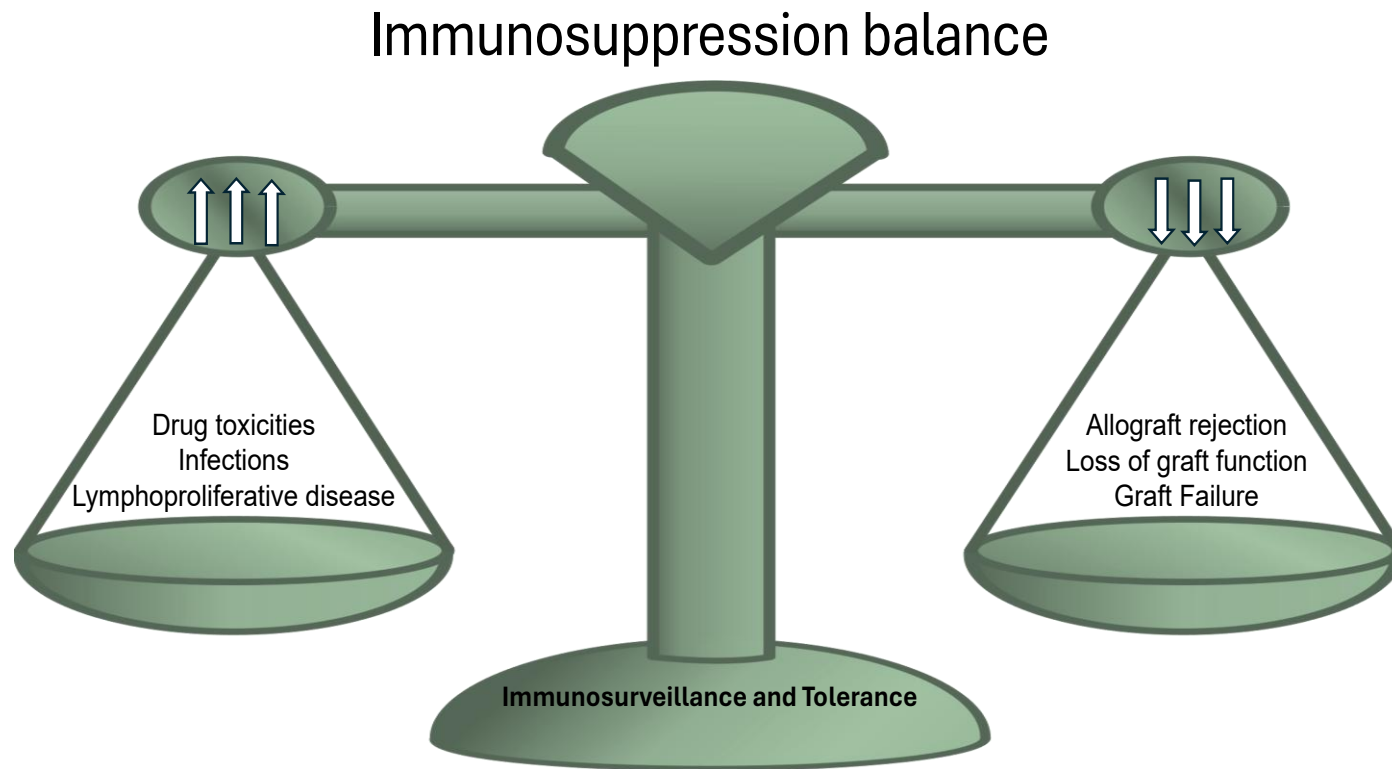
- Standardization of histological classification of rejection in biopsies (Banff Foundation for Allograft Pathology and ISHLT).
- Move from qualitative viral testing to quantitative (BK/CMV)- viral DNA-PCR.
- Dosing protocols/guidelines vary between transplant centers and the type of organ transplanted.
- More medications available as generic.
- More sizes/strengths/dosage forms of medications/extended-release formulations.

# Precision Medicine

Where we have been?

- “One size fits all” for immunosuppression: tacrolimus, mycophenolate and prednisone.
- Balance between overimmunosuppression and underimmunosuppression.
- We run immunosuppression the same regardless of underlying disease, age, race, or sex.
- Standard dosing based on weight and titrated to desired level.
- Immunosuppression is adjusted based on a goal to reduce and avoid toxicities while preventing rejection.
  - Organ specific tests (Creatine, Alk Phos, AST/ALT, WBC, PLT).

# Approaching Precision Immunosuppression Management



- Therapeutic Drug Monitoring
- Prospective Immunophenotyping to guide agent selection & dosing
- Risk stratification for adverse effects

### Neurological problems

- Tremors
- Cognitive slowing
- Headaches

### Cardiovascular disease

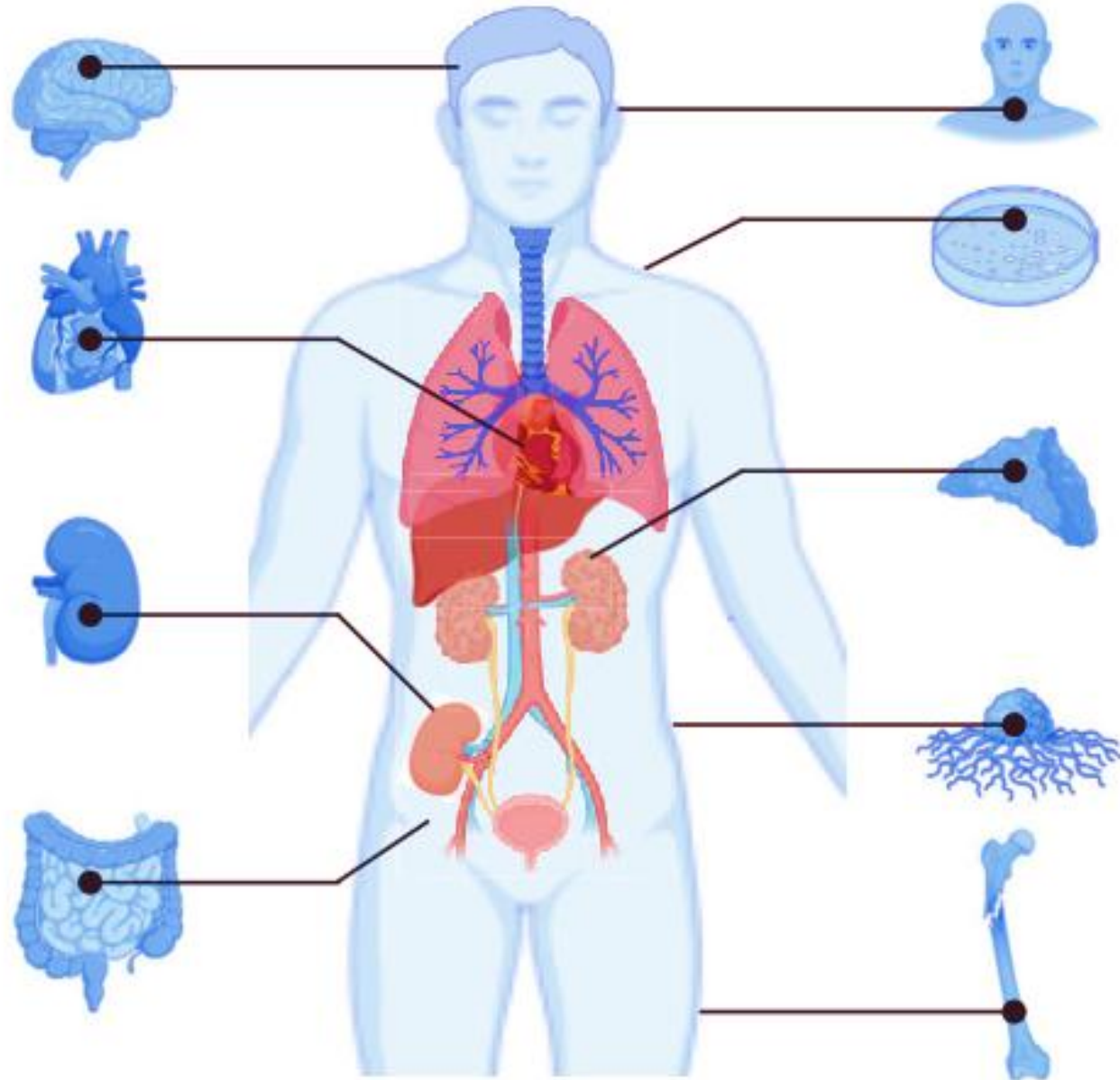
- Coronary artery disease
- Hypertension

### Kidney disease

- Reduced kidney filtration function
- Hyperkalaemia

### Gastrointestinal distress

- Nausea
- Diarrhoea



### Cosmetic side-effects

- Hair loss
- Gum hyperplasia

### Immune dysfunction

- Infections

### Endocrine disease

- Diabetes
- Hyperlipidaemia

### Cancer

- Skin
- Lymphoproliferative disorders
- Virally mediated malignancy

### Bone disease

- Fractures

# Precision Medicine

...and where are we headed?

- Immunosuppression is adjusted based on a goal to reduce and avoid toxicities while preventing rejection.
  - Organ specific tests (Creatine, Alk Phos, AST/ALT, WBC, PLT).
- Current advancements look at tailored immunosuppressive strategies.
- Understanding how genetic factors contribute to disease and using genetic markers to guide pharmacotherapy.
- Detecting disease at an early stage - From "reactive to proactive!".
- Minimizing invasive testing - "biopsy!"

# Challenges with Immunosuppression

- Tacrolimus has wide pharmacokinetic variability and inter-patient / intra-patient variability.
- Dose requirements dependent upon:
  - Age, Race, Sex, Body Weight, hepatic metabolism, and other co-administered drugs.
- Mycophenolic Acid: polymorphisms in UGT1A9 impact MPA clearance.
  - Unpredictable! Obtaining levels is difficult because of pharmacokinetic variability. No evidenced based recommendation for therapeutic range.
  - Dose adjustments are reactionary and based on clinical outcomes (GI side effects, neutropenia.)
  - Imuran: risk of myelosuppression based on genetics.
- Variability in drug pharmacokinetics can affect rejection risk.

# Challenges with Immunosuppression

- Age – variation in pediatric and elderly populations
- Race – genetic variation across different ethnic groups
- Short Telomere Syndrome (ILD)
- Patients with diseases that affect absorption/metabolism
  - Cystic Fibrosis
- Pre-transplant infections/colonization
- Patients with malignancies
- Patients with kidney disease
- Financial burden to patients
- Managing side effects and promoting compliance

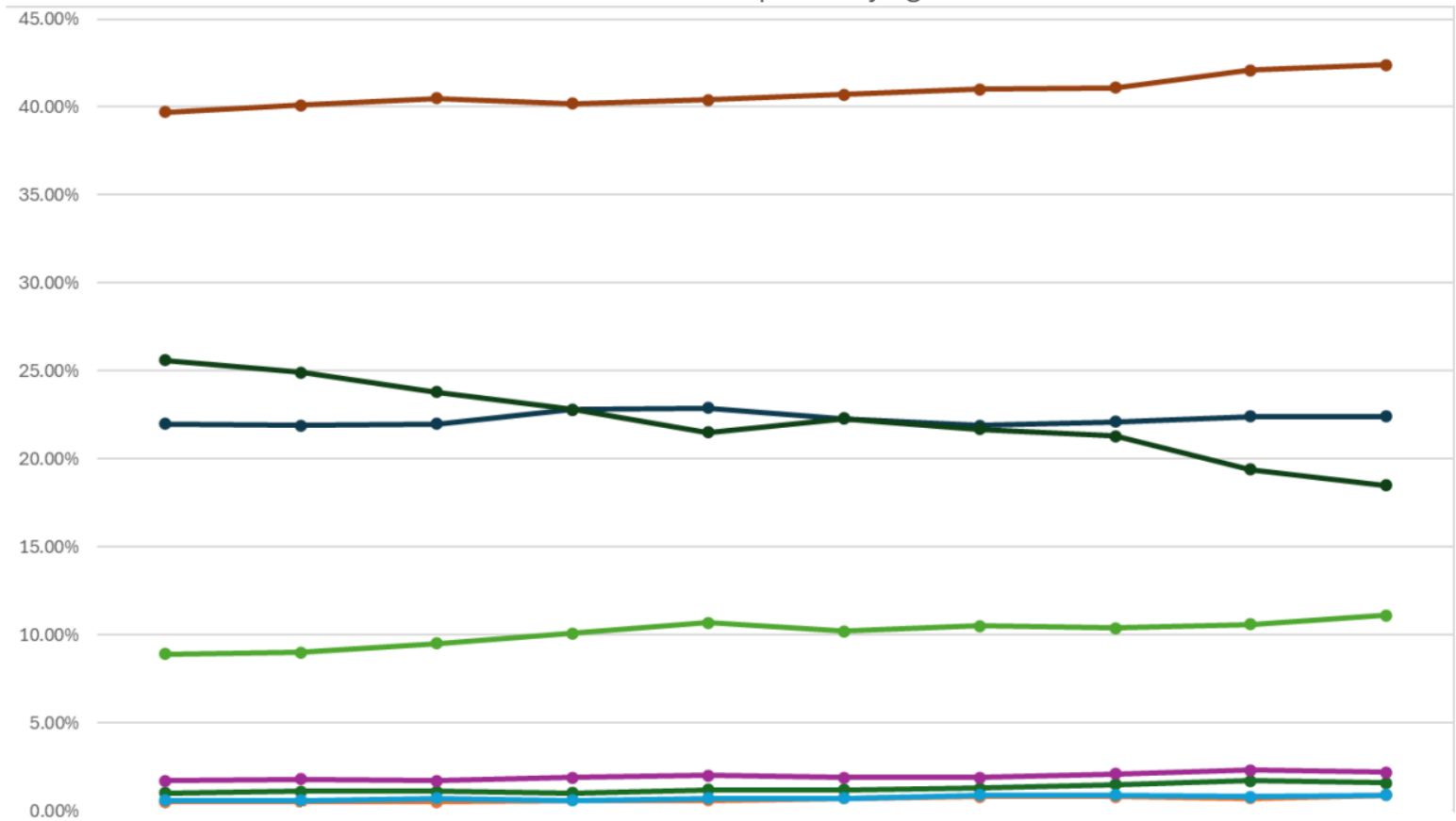
# Aging population and Transplant

- Rising aging population hence more elderly transplant recipients (>65 years).
- Improved short term graft survival and decreased acute rejection events.
- Higher risk of chronic allograft loss due to age related immune and nonimmune factors.
- Increased risk of infection, cardiovascular disease, malignancy, diabetes, fractures, myopathy ( higher risk of death with functioning graft).
- Decreased renal/hepatic clearance of drugs.
- Decrease cytochrome P<sub>450</sub> enzyme concentration.
- Reduce immunogenicity.

# Aging population and Transplant

- Retrospective Study looking at immunosuppression in elderly Kidney recipients: using US Transplant Registry data from 2005-2016 (14,887 patients  $\geq 65$ )
  - Lower risk of acute rejection 6 months-3 years post-transplant compared to younger patients with all immunosuppression regimens except IL2rAb + triple therapy.
  - Older adults are less likely to tolerate 3 drug immunosuppression.
    - Lower- intensity immunosuppression regimens including steroid sparing had statistically equivalent graft outcomes compared to T cell depleting + triple drug immunosuppression regimens.
  - CSA based, mTOR based, and Tacrolimus + anti-metabolite avoidance regimens were associated with poorer graft outcomes.
  - A reduced immunosuppressive regimen is recommended particularly steroid sparing (Tacrolimus/MMF/Steroid minimization or withdrawal).

National Transplants by Age



|               | 2025   | 2024   | 2023   | 2022   | 2021   | 2020   | 2019   | 2018   | 2017   | 2016   |
|---------------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|
| < 1 Year %    | 0.50%  | 0.50%  | 0.50%  | 0.60%  | 0.60%  | 0.70%  | 0.80%  | 0.80%  | 0.70%  | 0.90%  |
| 1-5 Years %   | 1.00%  | 1.10%  | 1.10%  | 1.00%  | 1.20%  | 1.20%  | 1.30%  | 1.50%  | 1.70%  | 1.60%  |
| 6-10 Years %  | 0.60%  | 0.60%  | 0.70%  | 0.60%  | 0.70%  | 0.70%  | 0.90%  | 0.90%  | 0.80%  | 0.90%  |
| 11-17 Years % | 1.70%  | 1.80%  | 1.70%  | 1.90%  | 2.00%  | 1.90%  | 1.90%  | 2.10%  | 2.30%  | 2.20%  |
| 18-34 Years % | 8.90%  | 9.00%  | 9.50%  | 10.10% | 10.70% | 10.20% | 10.50% | 10.40% | 10.60% | 11.10% |
| 35-49 Years % | 22.00% | 21.90% | 22.00% | 22.80% | 22.90% | 22.30% | 21.90% | 22.10% | 22.40% | 22.40% |
| 50-64 Years % | 39.70% | 40.10% | 40.50% | 40.20% | 40.40% | 40.70% | 41.00% | 41.10% | 42.10% | 42.40% |
| 65 + %        | 25.60% | 24.90% | 23.80% | 22.80% | 21.50% | 22.30% | 21.70% | 21.30% | 19.40% | 18.50% |

< 1 Year %    1-5 Years %    6-10 Years %    11-17 Years %    18-34 Years %    35-49 Years %    50-64 Years %    65 + %

# The Aging Immune System

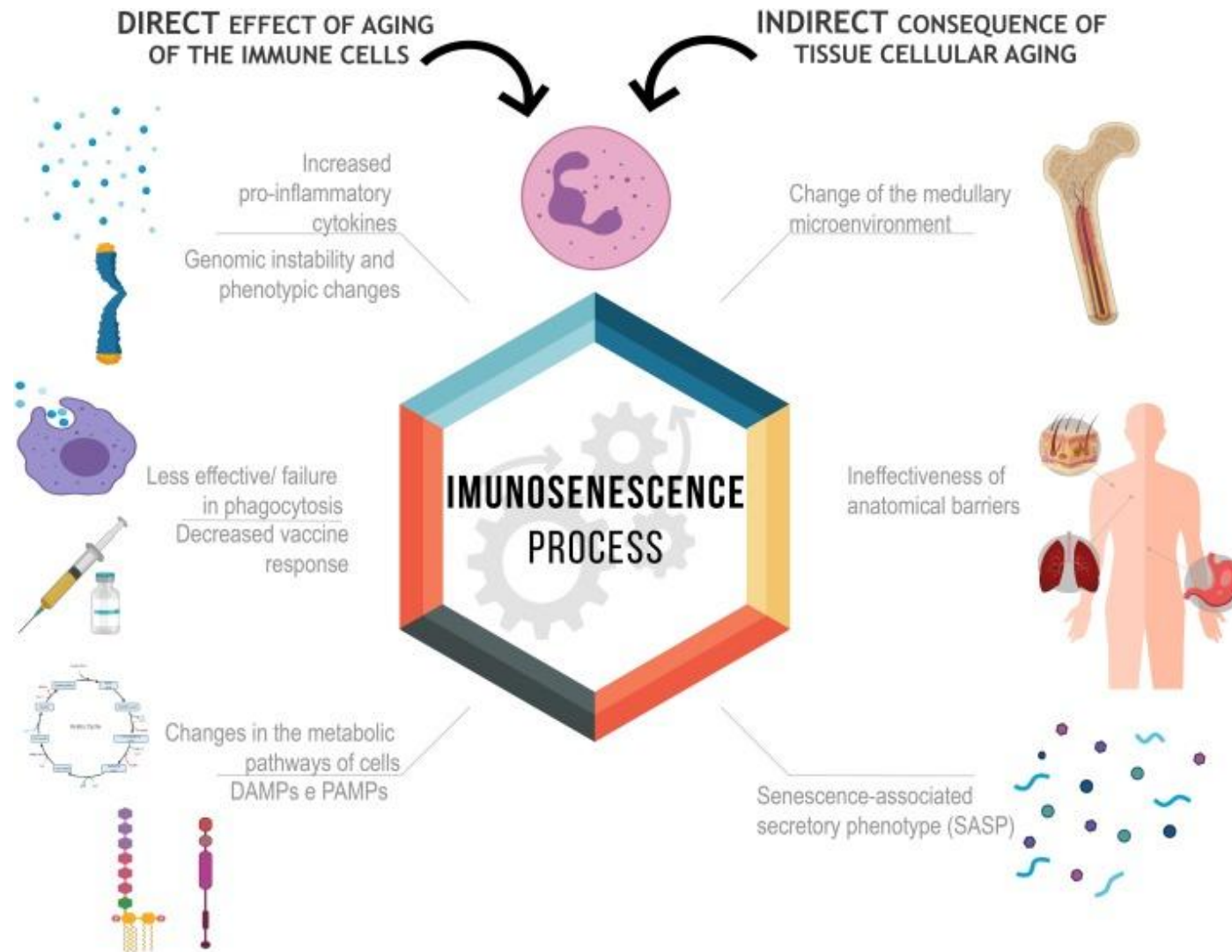
## "Immunosenescence"

- Decline in immune response
- Thymic involution, reduced naïve T and B cells, impaired antigen uptake/presentation,
- Impaired Wound Repair and decrease ability to fight infections/cancers, reduced vaccine responsiveness, coronary artery disease.

## "Cellular senescence" (Cell Cycle Arrest) and "Inflammaging"

- Mitochondrial dysfunction, decline in phagocytosis and shortened telomeres
- Senescent cells are not cleared as efficiently by the innate immune system.
  - Persistent low- grade inflammation (inflammaging)-increase in IL-6, TNF- $\alpha$ , CRP, IL-1 $\beta$
- Defective autophagy, activation of inflammasome and DNA damage response
  - Increase in senescent cells-stimulation of inflammasome-release of IL-1 $\beta$ /IL-18 (Senescent-Associated Secretory Phenotype)
- Loss of physical function, osteoarthritis, frailty, and loss of insulin sensitivity

# Cellular Senescence



- Aging → Defective Autophagy → Inflammasome Activation → DNA Damage → Persistent DDR → Senescence → SASP → Inflammation & Tissue Damage → More DNA Damage & Senescence → "Age Related Diseases"

## Precision Medicine and Transplant

- **What is it:** using biomarkers along with traditional clinical data to tailor and individualize medical care.
- Movement from reactive to proactive approach to care.
- Evaluating individual disease risk, evaluating for disease at an earlier stage, guiding drug dosing, guiding drug dosing, evaluating disease activity to estimate prognosis, and predict therapy responses.
- Precision health care incorporates genomics, patient and family history, comprehensive physical and clinical assessment, patient behaviors, and the patient's environment.
- Emergence of Targeted treatments.

# Types of Precision Testing

Pharmacogenetic testing

Donor derived Cell Free DNA

HLA DSA

Urinary mRNA (CXCL9, TIM-3)  
Reflecting renal T-cell/immune activation and predicts risk of rejection

High-resolution HLA-DQ/DP typing

BK/CMV viral load qPCR

CMV T cell immunity panels (CMV inSIGHT™ T Cell Immunity Panel)

KIR-ligand mismatch-May increase NK cell-mediated immune response to graft/rejection

Cytokines (TNF- $\alpha$ , IL-2, IFN- $\gamma$ )- Differences in the intensity of inflammatory responses regulated by cytokine polymorphisms

APOL1- high-risk genotype associated with graft functional decline

TLR4 rs49867-CMV/Bacterial Infection Susceptibility

AI Tools  
(diagnosing/prognosing/guiding therapy)

# Pharmacogenetics in Transplant

- Pharmacogenomics describes how one's genetic makeup affects drug metabolism
  - Correlates gene expression or single-nucleotide polymorphisms with drug efficacy and/or toxicity
- Clinical Pharmacogenetics Implementation Consortium (CPIC) provides evidence-based guidelines to interpret pharmacogenetic testing



# Pharmacogenetic Testing in Transplant

| Gene Tested                      | Medication   |
|----------------------------------|--------------|
| CYP <sub>3</sub> A <sub>5</sub>  | tacrolimus   |
| TPMT / NUDT <sub>15</sub>        | azathioprine |
| CYP <sub>2</sub> C <sub>19</sub> | voriconazole |
| G6PD                             | dapsone      |

# CYP<sub>3</sub>A<sub>5</sub>: tacrolimus

- Enzymes in the cytochrome P<sub>450</sub> (CYP) 3A family are responsible for the oxidative metabolism of tacrolimus.
  - CYP<sub>3</sub>A<sub>4</sub> and CYP<sub>3</sub>A<sub>5</sub> are thought to be relevant in adults
- First pass metabolism and systemic clearance of drugs metabolized by CYP<sub>3</sub>A<sub>5</sub> are susceptible to genetically determined differences in enzyme expression
- While CYP<sub>3</sub>A<sub>4</sub> poor metabolizers are rare, absence of functional CYP<sub>3</sub>A<sub>5</sub> is the norm in many populations.
  - most notable for white people
  - 80–85% of the population being homozygous for the variant CYP<sub>3</sub>A<sub>5</sub>\*<sub>3</sub> allele
- Retention of CYP<sub>3</sub>A<sub>5</sub> expression has been under some evolutionary selection pressure in populations originating close to the equator

# CYP<sub>3</sub>A<sub>5</sub>: tacrolimus

## CPIC GUIDELINES

**Table 1 Assignment of likely metabolism phenotypes based on CYP3A5 diplotypes**

| Likely phenotype                               | Genotypes                                                                    | Examples of diplotypes <sup>a</sup>      |
|------------------------------------------------|------------------------------------------------------------------------------|------------------------------------------|
| Extensive metabolizer<br>(CYP3A5 expresser)    | An individual carrying<br>two functional alleles                             | *1/*1                                    |
| Intermediate metabolizer<br>(CYP3A5 expresser) | An individual carrying one functional allele<br>and one nonfunctional allele | *1/*3, *1/*6, *1/*7                      |
| Poor metabolizer<br>(CYP3A5 nonexpresser)      | An individual carrying two nonfunctional alleles                             | *3/*3, *6/*6, *7/*7, *3/*6, *3/*7, *6/*7 |

<sup>a</sup>Additional rare variants, such as CYP3A5\*2, \*8, and \*9 may be found, which are of unknown functional significance. However, if a copy of \*1 is present, expected phenotype would be intermediate metabolizer.

# CYP3A5: tacrolimus

- “Poor metabolizer” generally indicates lower dose required
- ffor tacrolimus this indicates standard dose requirements

**Table 2 Dosing recommendations for tacrolimus based on CYP3A5 phenotype**

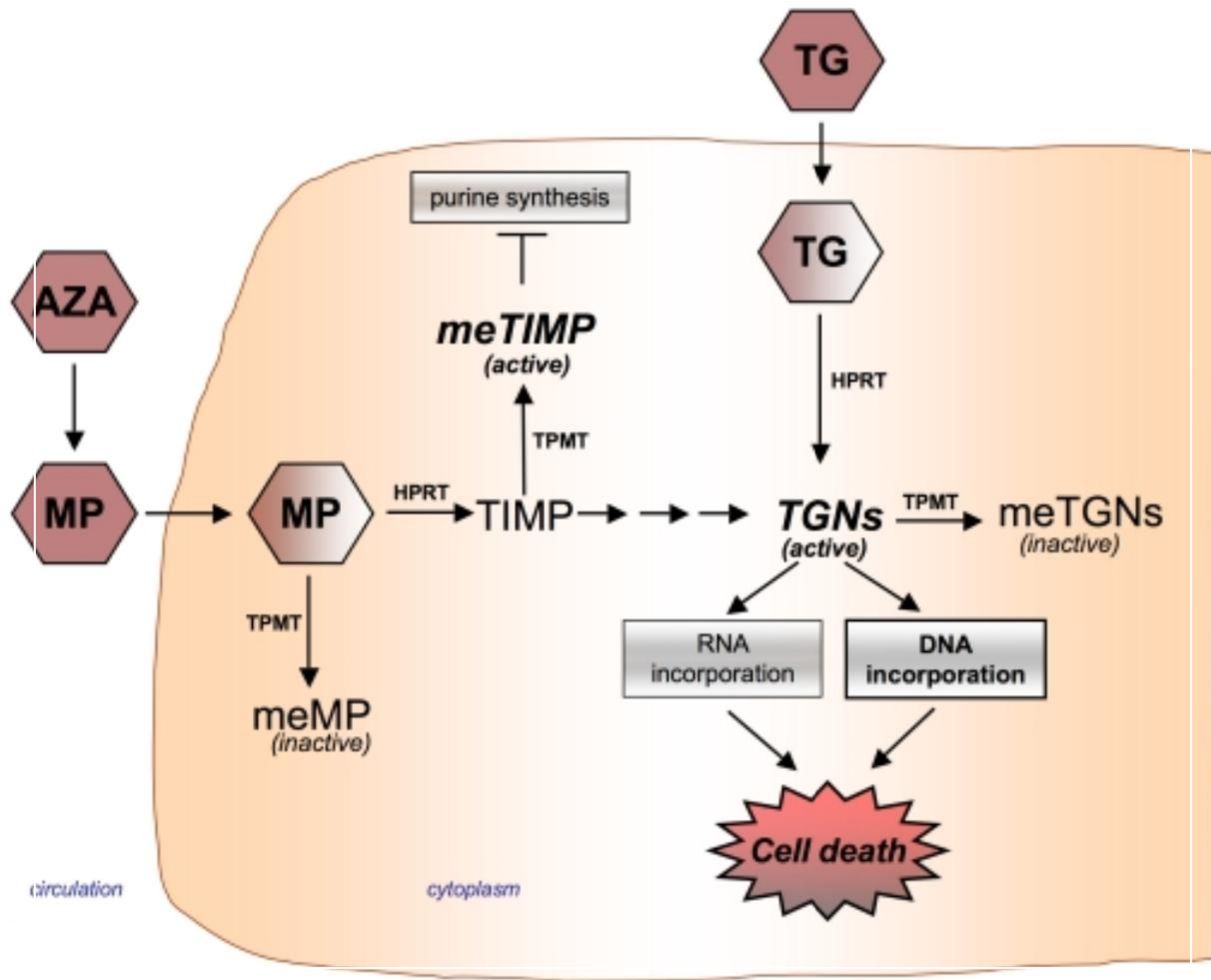
| CYP3A5 phenotype <sup>a</sup>               | Implications for tacrolimus pharmacologic measures                                                                                      | Therapeutic recommendations <sup>b</sup>                                                                                                                                                   | Classification of recommendations <sup>c</sup> |
|---------------------------------------------|-----------------------------------------------------------------------------------------------------------------------------------------|--------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|------------------------------------------------|
| Extensive metabolizer (CYP3A5 expresser)    | Lower dose-adjusted trough concentrations of tacrolimus and decreased chance of achieving target tacrolimus concentrations.             | Increase starting dose 1.5–2 times recommended starting dose. <sup>d</sup> Total starting dose should not exceed 0.3 mg/kg/day. Use therapeutic drug monitoring to guide dose adjustments. | Strong                                         |
| Intermediate metabolizer (CYP3A5 expresser) | Lower dose-adjusted trough concentrations of tacrolimus and decreased chance of achieving target tacrolimus concentrations.             | Increase starting dose 1.5–2 times recommended starting dose. <sup>a</sup> Total starting dose should not exceed 0.3 mg/kg/day. Use therapeutic drug monitoring to guide dose adjustments. | Strong                                         |
| Poor metabolizer (CYP3A5 nonexpresser)      | Higher (“normal”) dose-adjusted trough concentrations of tacrolimus and increased chance of achieving target tacrolimus concentrations. | Initiate therapy with standard recommended dose. Use therapeutic drug monitoring to guide dose adjustments.                                                                                | Strong                                         |

# CYP<sub>3</sub>A<sub>5</sub> and liver transplant

- In liver transplant recipients, the CYP<sub>3</sub>A<sub>5</sub> genotype of the donor liver may not be the same as the CYP<sub>3</sub>A<sub>5</sub> genotype of the recipient intestine.
- Often difficult to obtain donor pharmacogenomic testing
- Studies to date have been inconclusive as to the relative influence of the donor and recipient genotypes, and whether donor liver and recipient intestinal genotypes come into play at different points posttransplant.
  - some studies show that the donor genotype affects dose-adjusted trough concentrations from the first week posttransplant,
  - others show that it does not begin to play a role until the second week or even the sixth month posttransplant.

# TPMT / NUDT15: azathioprine

- Azathioprine is a thiopurine that is converted to mercaptopurine intracellularly
- TPMT is expressed in red blood cells, liver, and hematopoietic stem cells.
- TPMT catabolizes mercaptopurine to an inactive methylmercaptopurine base, leaving less parent drug available for potential anabolism to pharmacologically active forms
- Thioinosine monophosphate (intermediate metabolite of mercaptopurine) is also a substrate for TPMT
  - Downstream metabolites have pharmacologic activity: they inhibit de novo purine synthesis and may contribute to some of the adverse effects of mercaptopurine, particularly hepatotoxicity



**TPMT /  
NUDT15:  
AZATHIOPRINE**

# TPMT / NUDT15: azathioprine

**Table 1 Assignment of predicted TPMT and NUDT15 phenotypes based on genotypes**

| Predicted phenotype                                                 | Genotypes                                                                                                                                                                                                                      | Examples of diplotypes <sup>a</sup>                 |
|---------------------------------------------------------------------|--------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|-----------------------------------------------------|
| <i>Assignment of predicted TPMT phenotypes based on genotypes</i>   |                                                                                                                                                                                                                                |                                                     |
| Normal metabolizer                                                  | An individual carrying two normal function alleles<br>OR<br>one normal function allele PLUS one decreased function allele                                                                                                      | *1/*1<br>*1/*8                                      |
| Intermediate metabolizer                                            | An individual carrying one normal function allele PLUS one no-function allele<br>OR<br>two decreased function alleles                                                                                                          | *1/*2, *1/*3A, *1/*3B,<br>*1/*3C, *1/*4<br>*8/*8    |
| Possible intermediate metabolizer                                   | An individual carrying one uncertain/unknown function allele PLUS one no-function allele<br>OR<br>one no-function allele PLUS one decreased function allele                                                                    | *2/*9, *3A/*12<br>*3A/*8                            |
| Poor metabolizer                                                    | An individual carrying two no-function alleles                                                                                                                                                                                 | *3A/*3A, *2/*3A, *3A/*3C,<br>*3C/*4, *2/*3C, *3A/*4 |
| Indeterminate                                                       | An individual carrying one normal function allele PLUS one uncertain/unknown function allele<br>OR<br>one decreased function allele PLUS one uncertain/unknown function allele<br>OR<br>two uncertain/unknown function alleles | *1/*19<br>*8/*30<br>*19/*40                         |
| <i>Assignment of predicted NUDT15 phenotypes based on genotypes</i> |                                                                                                                                                                                                                                |                                                     |
| Normal metabolizer                                                  | An individual carrying two normal function alleles<br>OR<br>one normal function allele PLUS one decreased function allele                                                                                                      | *1/*1<br>*1/*5                                      |
| Intermediate metabolizer                                            | An individual carrying one normal function allele PLUS one no-function allele<br>OR<br>an individual carrying two decreased function alleles                                                                                   | *1/*2, *1/*3<br>*5/*5                               |
| Possible intermediate metabolizer                                   | An individual carrying one uncertain/unknown function allele PLUS one no-function allele                                                                                                                                       | *2/*15, *3/*21                                      |
| Poor metabolizer                                                    | An individual carrying two no-function alleles<br>OR<br>one no-function allele PLUS one decreased function allele                                                                                                              | *2/*2, *2/*3, *2/*4<br>*3/*5                        |
| Indeterminate                                                       | An individual carrying one normal function allele PLUS one uncertain/unknown function allele<br>OR<br>one decreased function allele PLUS one uncertain/unknown function allele<br>OR<br>two uncertain/unknown function alleles | *1/*12<br>*5/*20<br>*11/*14                         |

# TPMT / NUDT15: azathioprine

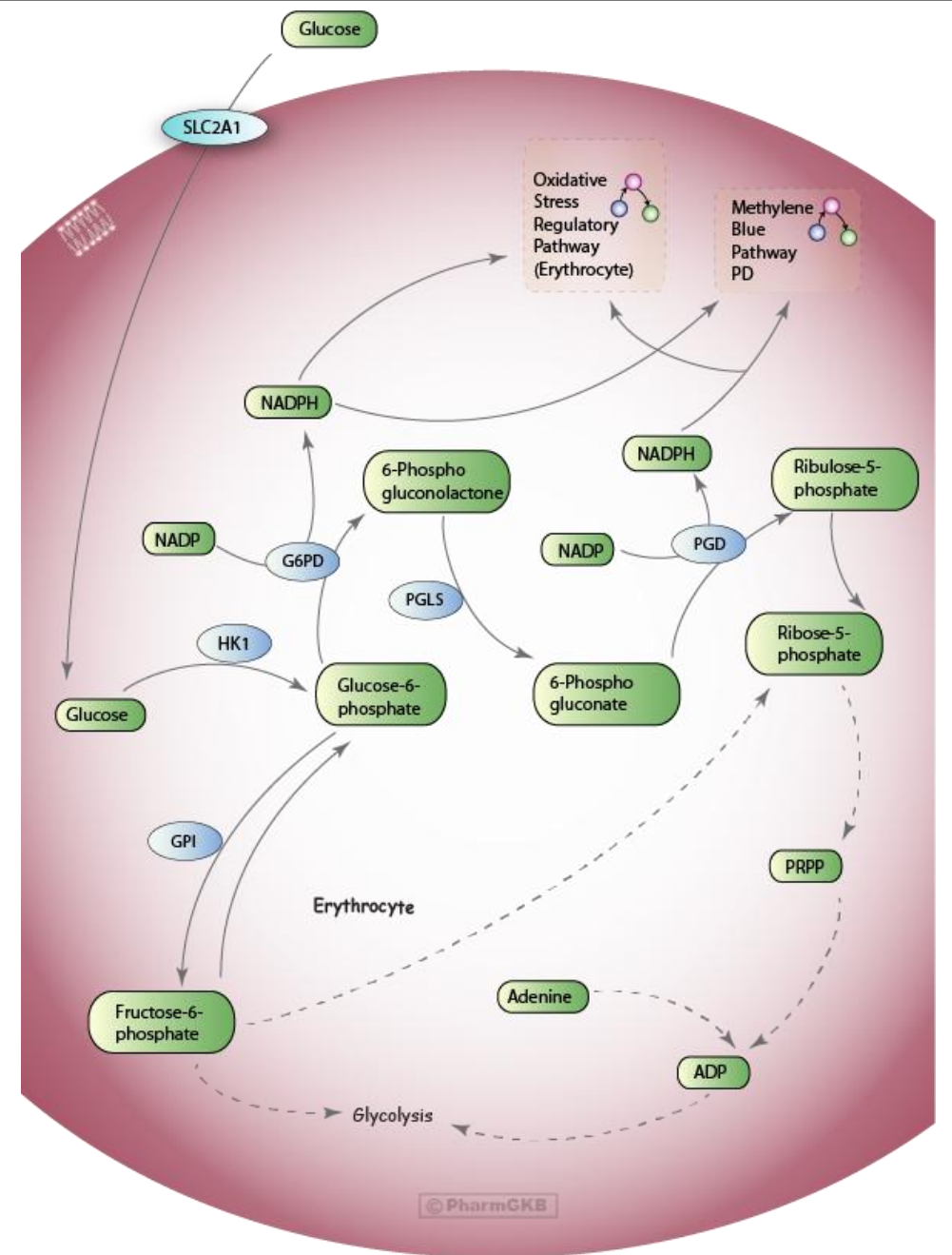
- A high dose of azathioprine (2mg/kg) exposes TPMT and/or NUDT15 poor metabolizers to extremely high concentrations of TGNs and potentially fatal myelosuppression.

**Table 4 Azathioprine dosing recommendations based on TPMT and/or NUDT15 phenotypes for nonmalignant conditions**

| TPMT/NUDT15 phenotype                                                                                                                               | Implications for toxicity                                                                                                                       | Implications for phenotypic measures <sup>a</sup>                                                                                                                      | Therapeutic recommendations <sup>b</sup>                                                                                                                                                                                                                                                                                                                                                                                                                                          | Classification of recommendation <sup>c</sup> |
|-----------------------------------------------------------------------------------------------------------------------------------------------------|-------------------------------------------------------------------------------------------------------------------------------------------------|------------------------------------------------------------------------------------------------------------------------------------------------------------------------|-----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|-----------------------------------------------|
| TPMT and NUDT15 normal metabolizer                                                                                                                  | Normal risk of thiopurine-related leukopenia, neutropenia and myelosuppression                                                                  | TPMT NMs have lower erythrocyte concentrations of TGN metabolites and higher concentrations of MeMPNs compared to TPMT IMs and TPMT PMs. This is the 'normal' pattern. | Initiate therapy with standard starting dose (e.g., 2 mg/kg/day for autoimmune diseases). During therapy, adjust doses of azathioprine based on disease-specific guidelines. It usually takes at least 2 weeks to reach steady state after each dose adjustment.                                                                                                                                                                                                                  | Strong                                        |
| TPMT normal metabolizer and NUDT15 (possible) intermediate metabolizer OR<br>TPMT (possible) intermediate metabolizer and NUDT15 normal metabolizer | Increased risk of thiopurine-related leukopenia, neutropenia and myelosuppression                                                               | TPMT IMs have moderate to high erythrocyte concentrations of TGN metabolites and low concentrations of MeMPNs compared to TPMT NMs.                                    | Initiate therapy with reduced starting doses (30–80% of standard starting dose) if standard starting dose is $\geq 2$ mg/kg/day<br>If starting dose is already below standard starting dose, dose reduction might not be necessary. During therapy, adjust the doses of azathioprine based on the degree of myelosuppression and disease-specific guidelines. It usually takes at least 2–4 weeks of stable dosing to reach steady state after each dose adjustment.              | Strong                                        |
| Any combination with TPMT poor metabolizer and/or NUDT15 poor metabolizer <sup>d</sup>                                                              | Greatly increased risk of thiopurine-related leukopenia, neutropenia and myelosuppression. <i>Fatal toxicity possible without dose decrease</i> | TPMT PMs have extremely high erythrocyte concentrations of TGN metabolites and no MeMPN compared to TPMT NMs.                                                          | Consider alternative nonthiopurine immunosuppressant therapy                                                                                                                                                                                                                                                                                                                                                                                                                      | Strong                                        |
| TPMT intermediate metabolizer and NUDT15 intermediate metabolizer (i.e., TPMT/NUDT15 compound intermediate metabolizer)                             | Increased risk of thiopurine-related leukopenia, neutropenia and myelosuppression<br>Higher risk of toxicity compared to single IM              | TPMT IMs have moderate to high erythrocyte concentrations of TGN metabolites and low concentrations of MeMPNs compared to TPMT NMs.                                    | Initiate therapy with reduced starting doses (20–50% of standard starting dose) if standard starting dose <sup>a</sup> is $\geq 2$ mg/kg/day<br>If starting dose is already below standard starting dose, dose reduction might not be necessary. During therapy, adjust the doses of azathioprine based on the degree of myelosuppression and disease-specific guidelines. It usually takes at least 2–4 weeks of stable dosing to reach steady state after each dose adjustment. | Moderate                                      |

# G6PD: dapsone

- The G6PD gene encodes the G6PD enzyme, which converts glucose-6-phosphate into 6-phosphogluconolactone, the first step of the pentose phosphate pathway



# G6PD: dapstone

## CPIC GUIDELINE

**Table 1 Assignment of predicted G6PD phenotype based on genotype**

| Predicted phenotype   | Genotype <sup>a</sup>                                                                                                                                                                    | Examples of G6PD genotypes <sup>b</sup>                                                                                                                                                                                 |
|-----------------------|------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|-------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| Normal                | A person with one X chromosome carrying a nondeficient (class IV) allele<br>A person carrying two nondeficient (class IV) alleles                                                        | B, Sao Borja, IV<br>B/B, B/Sao Borja, B/A, IV/IV                                                                                                                                                                        |
| Deficient             | A person with one X chromosome carrying a deficient (class II–III) allele<br>A person carrying two deficient (class II–III) alleles OR one class I allele and one class II or III allele | A-, Orissa, Kalyan-Kerala, Mediterranean, Canton, Chatham, II, III<br>A-/A-, A-/Orissa, Orissa/Kalyan-Kerala, Mediterranean/Mediterranean, Chatham/Mediterranean, Canton/Viangchan, II/II, II/III, III/III, I/II, I/III |
| Deficient with CNSHA  | A person with one X chromosome carrying a deficient (class I) allele<br>A person carrying two deficient (class I) alleles <sup>c</sup>                                                   | Bangkok, Villeurbanne, I<br>Bangkok/Bangkok, Bangkok/Villeurbanne, I/I                                                                                                                                                  |
| Variable <sup>d</sup> | A person carrying one nondeficient (class IV) allele and one deficient (class I–III) allele                                                                                              | B/Bangkok, B/Mediterranean, B/A-, IV/I, IV/II, IV/III                                                                                                                                                                   |
| Indeterminate         | A person carrying at least one uncertain function allele                                                                                                                                 | Laibin, Yunan<br>B/Yunan, II/Yunan, III/Laibin, IV/Yunan                                                                                                                                                                |

| Predicted G6PD phenotype based on genotype | Implications for phenotypic measures                 | Therapeutic recommendations for high-risk drugs                                                                                                     |
|--------------------------------------------|------------------------------------------------------|-----------------------------------------------------------------------------------------------------------------------------------------------------|
| Normal                                     | Low risk of acute hemolytic anemia                   | No reason to avoid high-risk drugs based on G6PD status                                                                                             |
| Deficient                                  | High risk of acute hemolytic anemia                  | Avoid use of high-risk drugs                                                                                                                        |
| Deficient with CNSHA                       | High risk of acute exacerbation of chronic hemolysis | Avoid use of high-risk drugs                                                                                                                        |
| Variable                                   | Variable risk of acute hemolytic anemia              | To ascertain G6PD status, enzyme activity must be measured. Drug use should be guided per the recommendations based on the activity-based phenotype |
| Indeterminate                              | Unknown risk of acute hemolytic anemia               | To ascertain G6PD status, enzyme activity must be measured. Drug use should be guided per the recommendations based on the activity-based phenotype |

# G6PD: dapstone

# CYP2C19: voriconazole

- Voriconazole demonstrates wide interpatient variability in serum concentrations, due in part to variant CYP2C19 alleles
- CYP2C19 is highly polymorphic with 35 defined variant star (\*) alleles

**Table 1 Assignment of likely CYP2C19 phenotypes based on genotypes**

| Likely phenotype                                                           | Genotypes <sup>a</sup>                                                                                                                   | Examples of CYP2C19 diplotypes    |
|----------------------------------------------------------------------------|------------------------------------------------------------------------------------------------------------------------------------------|-----------------------------------|
| CYP2C19 ultrarapid metabolizer (~2–5% of patients) <sup>b</sup>            | An individual carrying two increased function alleles                                                                                    | *17/*17                           |
| CYP2C19 rapid metabolizer (~2–30% of patients) <sup>b</sup>                | An individual carrying one normal function allele and one increased function allele                                                      | *1/*17                            |
| CYP2C19 normal metabolizer <sup>c</sup> (~35–50% of patients) <sup>b</sup> | An individual carrying two normal function alleles                                                                                       | *1/*1                             |
| CYP2C19 intermediate metabolizer (~18–45% of patients) <sup>b</sup>        | An individual carrying one normal function allele and one no function allele or one no function allele and one increased function allele | *1/*2, *1/*3, *2/*17 <sup>d</sup> |
| CYP2C19 poor metabolizer (~2–15% of patients) <sup>b</sup>                 | An individual carrying two no function alleles                                                                                           | *2/*2, *2/*3, *3/*3               |

# CYP<sub>2</sub>C<sub>19</sub>: voriconazole

- Alternative agents recommended for ultrarapid, rapid, and poor metabolizers

**Table 2 Dosing recommendations for voriconazole treatment based on CYP2C19 phenotype for adult patients**

| CYP2C19 phenotype                        | Implications for voriconazole pharmacologic measures                                                                                                                                    | Therapeutic recommendations                                                                                                                                                                                                                                                                                                                                                                                                                                                                     | Classification of recommendations <sup>a</sup> |
|------------------------------------------|-----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|-------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|------------------------------------------------|
| CYP2C19 ultrarapid metabolizer (*17/*17) | In patients for whom an ultrarapid metabolizer genotype (*17/*17) is identified, the probability of attainment of therapeutic voriconazole concentrations is small with standard dosing | Choose an alternative agent that is not dependent on CYP2C19 metabolism as primary therapy in lieu of voriconazole. Such agents include isavuconazole, liposomal amphotericin B, and posaconazole. <sup>b</sup>                                                                                                                                                                                                                                                                                 | Moderate <sup>c</sup>                          |
| CYP2C19 rapid metabolizer (*1/*17)       | In patients for whom a rapid metabolizer genotype (*1/*17) is identified, the probability of attainment of therapeutic concentrations is modest with standard dosing                    | Choose an alternative agent that is not dependent on CYP2C19 metabolism as primary therapy in lieu of voriconazole. Such agents include isavuconazole, liposomal amphotericin B, and posaconazole. <sup>b</sup>                                                                                                                                                                                                                                                                                 | Moderate                                       |
| CYP2C19 normal metabolizer               | Normal voriconazole metabolism                                                                                                                                                          | Initiate therapy with recommended standard of care dosing <sup>b</sup>                                                                                                                                                                                                                                                                                                                                                                                                                          | Strong                                         |
| CYP2C19 intermediate metabolizer         | Higher dose-adjusted trough concentrations of voriconazole compared with normal metabolizers                                                                                            | Initiate therapy with recommended standard of care dosing <sup>b</sup>                                                                                                                                                                                                                                                                                                                                                                                                                          | Moderate                                       |
| CYP2C19 poor metabolizer                 | Higher dose-adjusted trough concentrations of voriconazole and may increase probability of adverse events                                                                               | Choose an alternative agent that is not dependent on CYP2C19 metabolism as primary therapy in lieu of voriconazole. Such agents include isavuconazole, liposomal amphotericin B, and posaconazole. <sup>b</sup> In the event that voriconazole is considered to be the most appropriate agent, based on clinical advice, for a patient with poor metabolizer genotype, voriconazole should be administered at a preferably lower than standard dosage with careful therapeutic drug monitoring. | Moderate                                       |

## Pharmacogenomic Results

→ CYP2C19 \*1/\*17 ( Drug response )

Genotypes: CYP2C19 \*1/\*17

CYP2C9 \*1/\*1 ( Drug response )

Genotypes: CYP2C9 \*1/\*1

CYP2D6 (\*4/\*4)2N ( Drug response )

Genotypes: CYP2D6 (\*4/\*4)2N

→ CYP3A5 \*3/\*3 ( Drug response )

Genotypes: CYP3A5 \*3/\*3

CYP4F2 \*1/\*3 ( Drug response )

Genotypes: CYP4F2 \*1/\*3

DPYD Reference/Reference ( Drug response )

Genotypes: DPYD Reference/Reference

→ NUDT15 \*1/\*1 ( Drug response )

Genotypes: NUDT15 \*1/\*1

SLCO1B1 \*1/\*1 ( Drug response )

Genotypes: SLCO1B1 \*1/\*1

→ TPMT \*1/\*1 ( Drug response )

Genotypes: TPMT \*1/\*1

VKORC1 c.-1639G>A/c.-1639G>A ( Drug response )

Genotypes: VKORC1 c.-1639G>A/c.-1639G>A

# Practical Application

# Practical Application

- CYP2C19 \*1/\*17
  - Voriconazole rapid metabolizer
- CYP3A5 \*3\*3
  - Tacrolimus poor metabolizer
- NUDT15 / TPMT \*1\*1
  - Azathioprine normal metabolizer

## Pharmacogenomic Results

→ CYP2C19 \*1/\*17 ( Drug response )

Genotypes: CYP2C19 \*1/\*17

CYP2C9 \*1/\*1 ( Drug response )

Genotypes: CYP2C9 \*1/\*1

CYP2D6 (\*4/\*4)2N ( Drug response )

Genotypes: CYP2D6 (\*4/\*4)2N

→ CYP3A5 \*3/\*3 ( Drug response )

Genotypes: CYP3A5 \*3/\*3

CYP4F2 \*1/\*3 ( Drug response )

Genotypes: CYP4F2 \*1/\*3

DPYD Reference/Reference ( Drug response )

Genotypes: DPYD Reference/Reference

→ NUDT15 \*1/\*1 ( Drug response )

Genotypes: NUDT15 \*1/\*1

SLCO1B1 \*1/\*1 ( Drug response )

Genotypes: SLCO1B1 \*1/\*1

→ TPMT \*1/\*1 ( Drug response )

Genotypes: TPMT \*1/\*1

VKORC1 c.-1639G>A/c.-1639G>A ( Drug response )

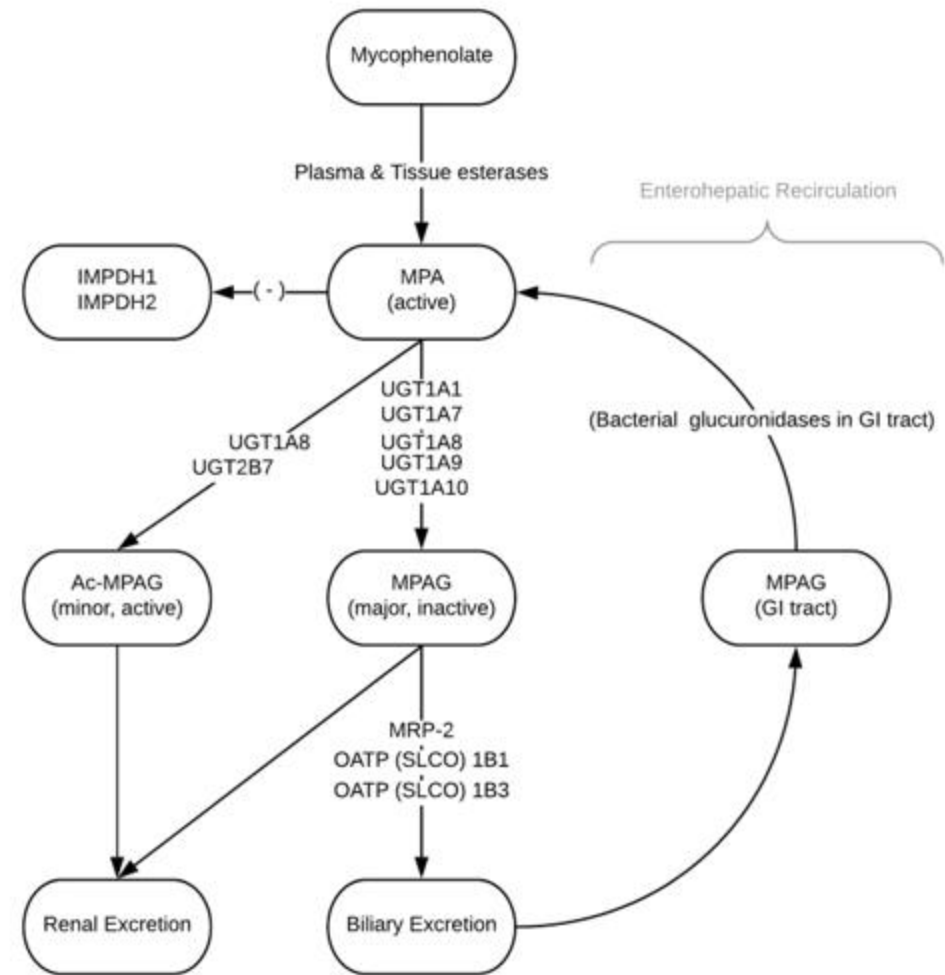
Genotypes: VKORC1 c.-1639G>A/c.-1639G>A

# UGT1A9: mycophenolate mofetil

- Mycophenolic acid (MPA), the active metabolite of the prodrug mycophenolate mofetil (MMF), is a standard immunosuppressive drug used after hematopoietic stem cell and solid organ transplantation.
- The pharmacokinetic profile of the drug and its phenolic (MPAG) and acyl (AcMPAG) glucuronides is characterized by unexplained interindividual variation.
- CPIC does not have a published clinical guideline for UGT1A9 and its impact on MMF.

# Mycophenolate therapeutic drug monitoring

- Single time point drug levels not accurate due to complicated pharmacokinetics
  - Non-linear absorption kinetics
  - Enterohepatic circulation
- Estimated MPA area under the curve (AUC) is more successful
  - Impractical because it requires multiple lab draws across several hours



# **ROLE OF THE APP IN PRECISION MEDICINE**

# Precision Medicine and the APP- Implementation

- Obtain Thorough History:
  - Family
  - Patient
- Clinical Assessment
- Assess Patient Behavior
- Economic Issues/Financial Burden
- Evaluate environmental factors
- Biomedical Results (labs, biopsies, PFT's, Echo)
- Review of Imaging/Ordering New Imaging
- Goal is to determine disease susceptibility, exposures, and potential response/lack of response to treatment
- Initiate precision testing pre-transplant/post-transplant

# Precision Medicine at all Phases of Transplant

- Pre-Transplant:
  - High Resolution HLA Typing
  - PRA (panel reactive antibody)
  - CYP3A5 Genotype/Polymorphisms
  - UGT1A9-Mycophenolic acid exposure/toxicity
  - TPMT and NUDT15 –azathioprine metabolism/risk of myelosuppression
- Peri-Operatively: Pharmacogenetic Guide Dosing
  - Refer to results of pharmacogenetic testing
- Post-Transplant: Non-invasive rejection Monitoring
  - dd-cfDNA
  - HLA DSA (donor specific anti-HLA antibodies)
  - CMV/BK Virus Monitoring

# Donor Derived Cell Free DNA (dd-cfDNA)

- DNA fragments released in bloodstream and detected at molecular level
  - Allosure-CareDx and Pospera-Natera
  - DNA fragments released during cellular apoptosis and necrosis
- Non-invasive biomarker of allograft injury(not necessarily specific to rejection but can be a marker of rejection). Can also be a marker of subclinical graft injury in the absence of clinical symptoms.
- Absolute dd-cfDNA values are not directly comparable across laboratories due to differences in extraction methods, sample preservation, and quantification techniques

# Donor Derived Cell Free DNA (dd-cfDNA)

- May be used in place of or in conjunction with a biopsy.
- May be used to direct/tailor immunosuppression.
- Combining dd-cfDNA with other biomarkers:
  - DSA testing
  - Labs
  - Imaging in addition to clinical picture can help guide clinical decisions
- Recommended to measure serial measurements and monitor trends (false-positives and false-negatives).
- Can be elevated in setting of infection, ischemia, or inflammation.
- Interpretation based on:
- Type of transplant
  - Time post-transplant
  - Clinical picture (malignancy, infection, leukopenia, ischemia).

# Donor Derived Cell Free DNA (dd-cfDNA) and Role of the APP

- Establish protocol for obtaining dd-cfDNA in the post-transplant recipient.
- Identify high risk patients that could benefit from dd-cfDNA vs biopsy.
- Educate patient as to usefulness of dd-cfDNA.
- Make sure samples are obtained in correct tube as provided by the company running the test (cannot use conventional EDTA tubes).
- Providers typically supply collection kits, prepaid shipping, and electronic reporting.
- Obtain dd-cfDNA along with DSA, biopsy, lab, imaging, and/or clinical evaluation to drive treatment or observation. Biopsy is still “Gold Standard”.
- Using dd-cfDNA may minimize the need for invasive procedure, guide treatment, and detect injury earlier.

# Cell Free DNA (dd-cfDNA)

## What is currently available?

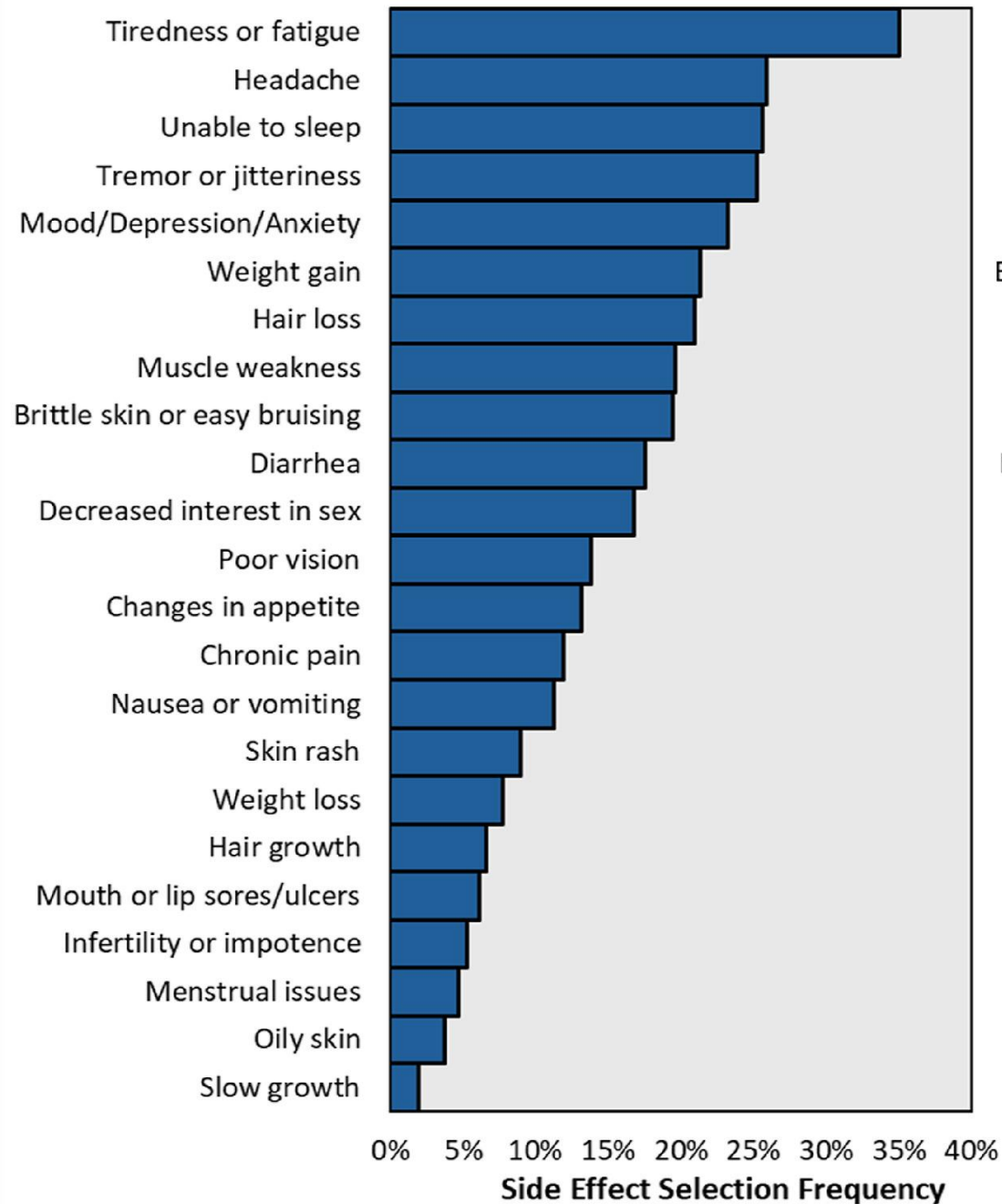
- **Allosure (Caredx)**
- **Prospera (Natera)**
- TRAC (Viracor Eurofins)
- AlloNext (Eurofins Genome)
- One Lambda Devyser Accept cfDNA (Devyser)
- GraftAssureCore (previously VitaGraft)
- iMDx (previously Oncocyte)
- HoloGRAFT ONE (Omixon)

# **WHAT DOES ALL THIS MEAN FOR PATIENTS?**

"The Patient Experience"

# Immunosuppression and Side Effects

- American Society of Transplantation: Web-based study of 10,091 responses included (9543 adults, 548 pediatric respondents) to assess burden of immunosuppression and quality of life.
- **92%** reported at least one side effect and **63%** reported side effect as occurring "often" or "always"
- **54%** of side effects were rated as having a "moderate" or "great deal" of impact on daily life.
- Those  $\geq 60$  years had **23%** higher side effect burden compared to  $< 60$  year
- Immunosuppression causes side effects and heavy burden to patients.
- More concern among respondents about infection and cancer versus graft loss and rejection although still rated as high concern.
- Most concerning issues related to self-care: interference related to travel for work or vacation, feeling dependent, interference in hobbies or daily activities.
- Issues with lowest concern: tracking medical appointments, keeping appointments, monitoring health conditions.
- Despite burden adherence is HIGH. (**25%** skipped doses due to side effects and **40%** missed refill due to cost)





# Precision Medicine & the APP

# Precision Medicine and the Advanced Provider

- Personalize and tailor care to improve care while maintaining a holistic approach to care.
- Implement biomarker testing.
- Develop a treatment plan.
- Order specific tests and/or biopsies.
- Administer therapy and monitor/manage adverse effects.
- Collaborate with pharmacist and physician to coordinate care (multidisciplinary).
- Educate patients and colleagues about precision testing.
- Aim to improve graft survival, reduce cost due to lower immunosuppression, and avoid side effects.

# Summary

- Immunosuppression continues to evolve and we are trying to move from a "one size fits all" approach to a more patient specific and tailored strategy.
- Certain patient populations present challenges to immunosuppression, and this must be taken into consideration.
- In the last 10-20 years there has been an increase in transplant recipients 65 years and older and precision medicine could lend insight into how this patient population's more fragile immune system is impacted by immunosuppression.
- Despite side effects of immunosuppression there is high compliance however, precision medicine offers a better balance between over and underimmunosuppression while minimizing side effects.
- Precision medicine should be implemented at specific times in the transplant phase to help guide treatment and improve outcomes.
- Recommended precision testing if feasible: pharmacogenetic testing, HLA DSA, dd-cfDNA, Viral PCR testing to name a few can be implemented.
- The APP is an integral part of the interdisciplinary team who can determine how and when to implement precision testing in the transplant recipient.
- It is crucial for the APP to be aware of what precision testing is available and educate the patient and colleagues about these tests.
- Leverage patient outcomes, treatment burden, and graft survival.