

# **Personalized Antiplatelet Therapy Possible? Not so fast...**

Angioplasty Summit – TCAP Meeting 2011

Guillaume Pare MD

Canada Research Chair in Genetic and Molecular Epidemiology



# Disclosures

- I have received research support and honoraria from BMS and sanofi-aventis
- As a geneticist, I strongly believe routine personalized medicine based on genotypes will eventually be an important component of patient care!

# Outline

- Using the right outcome to assess efficacy and safety of antiplatelet therapy
- Does CYP2C19/platelet function testing (PFT) provide clinically significant prognostic value?
- Does intervention that alters the risk factor lead to clinical benefit?

# Choosing the right outcome

- Neither the effect of CYP2C19 genotypes/PFT nor the causality of the association are at stake in PCI patients



- The effect of CYP2C19 genotypes and PFT varies according to outcome
  - Stent thrombosis seems particularly sensitive

# Choosing the right outcome

- Any outcome really is a surrogate for mortality and patient well-being
- Patient outcome should be as representative of these difficult to measure endpoints
  - Need to be inclusive in our definition
- Major Adverse Cardiovascular Events best captures all types of events but...

# Choosing the right outcome



MACE (11.7%)  
MI (6.9%)  
Death from vascular causes (5.1%)  
Stent thrombosis (1.9%; definite)  
Ischemic stroke (1.1%)

Major bleeding (11.2%)  
Life-threatening or fatal bleeding (5.8%)  
Fatal bleeding (0.3%)

Taken from the PLATO study, clopidogrel arm  
Wallentin et al. NEJM 361(11) 2009: 1045-57

# Does CYP2C19 testing/PFT provide clinically significant prognostic value?

- P-values alone are not representative of clinical usefulness
- Effect size matters!
  - HR of 3.0 not the same as 1.5
- Highest ORs tend to be reported by smaller studies

# Does CYP2C19 testing/PFT provide clinically significant prognostic value?

- CYP2C19 effect on MACE in studies >500 individuals

| Study                   | %PCI      | Loss-of-function Carriers |             | No Loss-of-Function Allele |              | Unadj. OR (95%CI)       | p           |
|-------------------------|-----------|---------------------------|-------------|----------------------------|--------------|-------------------------|-------------|
|                         |           | No. Events                | No. at Risk | No. Events                 | No. at Risk  |                         |             |
| EXCELSIOR               | 100       | 5                         | 243         | 7                          | 554          | 1.65 (0.47-5.36)        | 0.53        |
| TRITON                  | 100       | 46                        | 395         | 83                         | 1064         | 1.56 (1.06-2.27)        | 0.03        |
| FAST-MI                 | 70        | 63                        | 635         | 193                        | 1573         | 0.79 (0.58-1.06)        | 0.12        |
| RECLOSE                 | 100       | 15                        | 247         | 14                         | 525          | 2.36 (1.11-5.05)        | 0.03        |
| ISAR                    | 100       | 55                        | 680         | 119                        | 1805         | 1.25 (0.89-1.73)        | 0.22        |
| Intermountain           | 100       | 68                        | 344         | 141                        | 906          | 1.34 (0.97-1.84)        | 0.09        |
| PLATO                   | 66        | 149                       | 1388        | 332                        | 3516         | 1.15 (0.94-1.41)        | 0.18        |
| CURE                    | 15        | 52                        | 651         | 179                        | 1886         | 0.83 (0.60-1.14)        | 0.27        |
| <b>OVERALL</b>          | <b>73</b> | <b>453</b>                | <b>4583</b> | <b>1068</b>                | <b>11829</b> | <b>1.12 (0.99-1.26)</b> | <b>0.06</b> |
| <b>OVERALL w/o CURE</b> | <b>83</b> | <b>401</b>                | <b>3932</b> | <b>889</b>                 | <b>9943</b>  | <b>1.17 (1.04-1.33)</b> | <b>0.01</b> |

Unadjusted analysis is relevant since genotypes are unlikely to be adjusted for patient characteristics to determine therapy



# Does CYP2C19 testing/PFT provide clinically significant prognostic value?

- Comparing “poor metabolizers” to individuals without any LOF in studies >500 participants

| Study                   | %PCI      | Two Loss-of-function Alleles |             | No Loss-of-Function Allele |             | Unadj. OR (95%CI)       | p           |
|-------------------------|-----------|------------------------------|-------------|----------------------------|-------------|-------------------------|-------------|
|                         |           | No. Events                   | No. at Risk | No. Events                 | No. at Risk |                         |             |
| TRITON                  | 100       | 4                            | 38          | 83                         | 1064        | 1.44 (0.41-3.73)        | 0.53        |
| FAST-MI                 | 70        | 10                           | 58          | 193                        | 1573        | 1.51 (0.71-2.91)        | 0.31        |
| RECLOSE                 | 100       | 2                            | 26          | 14                         | 525         | 3.21 (0.44-12.6)        | 0.17        |
| ISAR                    | 100       | 3                            | 47          | 119                        | 1805        | 1.01 (0.23-2.83)        | 1           |
| Intermountain           | 100       | 3                            | 14          | 141                        | 906         | 1.53 (0.33-5.07)        | 0.47        |
| CURE                    | 15        | 4                            | 61          | 178                        | 1880        | 0.70 (0.20-1.72)        | 0.65        |
| <b>OVERALL</b>          | <b>73</b> | <b>26</b>                    | <b>244</b>  | <b>728</b>                 | <b>7753</b> | <b>1.22 (0.80-1.85)</b> | <b>0.36</b> |
| <b>OVERALL w/o CURE</b> | <b>91</b> | <b>22</b>                    | <b>183</b>  | <b>550</b>                 | <b>5873</b> | <b>1.43 (0.90-2.27)</b> | <b>0.13</b> |

# Does CYP2C19 testing/PFT provide clinically significant prognostic value?

- Similar results for PFT
  - Smaller studies report higher ORs (>6 in some cases)
  - Compounded by lack of predetermined threshold
  - HR 1.68 (95%CI 0.76-3.72; p=0.2) in GRAVITAS
  - Further complicated by entanglement of pharmacologic effect and intrinsic platelet factors on platelet reactivity

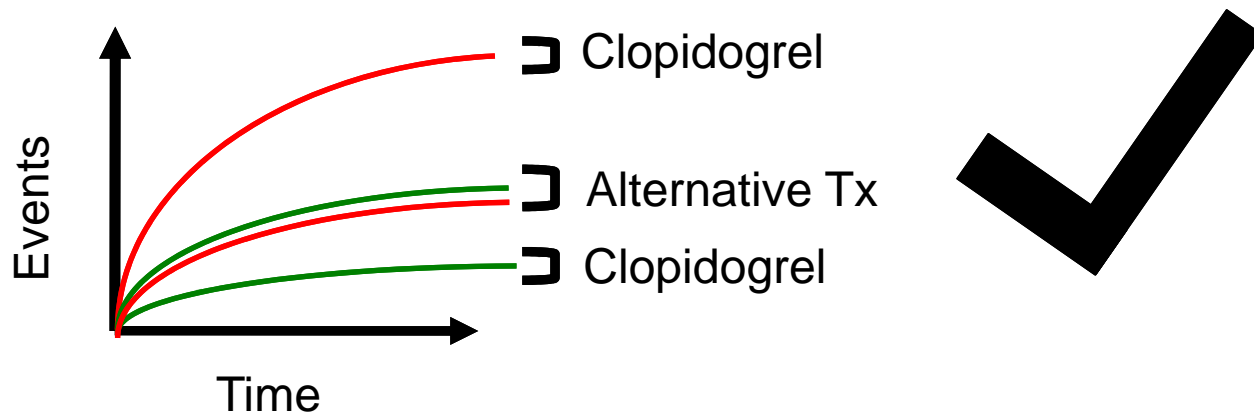
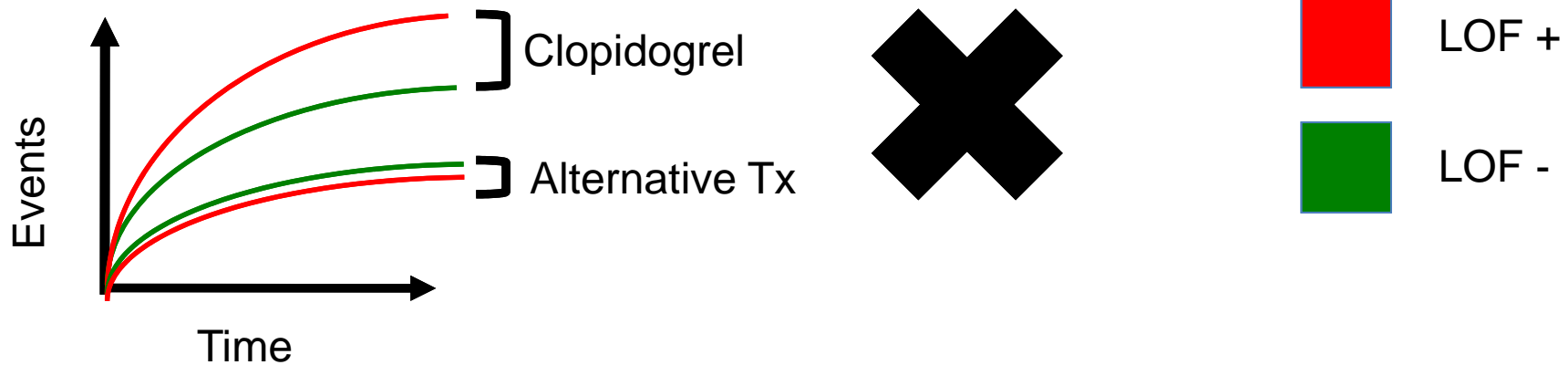
Bonello et al. JACC 56(12) 2010: 919-33

Price et al. JAMA 305(11) 2011: 1097-1105

# Does intervention that alters the risk factor lead to clinical benefit?

- Association of CYP2C19 genotypes with outcome is not sufficient to warrant clinical testing
- To justify testing, there needs to be clear evidence that either:
  - A group of patients will derive greater benefit from clopidogrel than the alternative antiplatelet treatment
  - A group of patient will derive equal benefit from clopidogrel as the more expensive alternative

# Does intervention that alters the risk factor lead to clinical benefit?



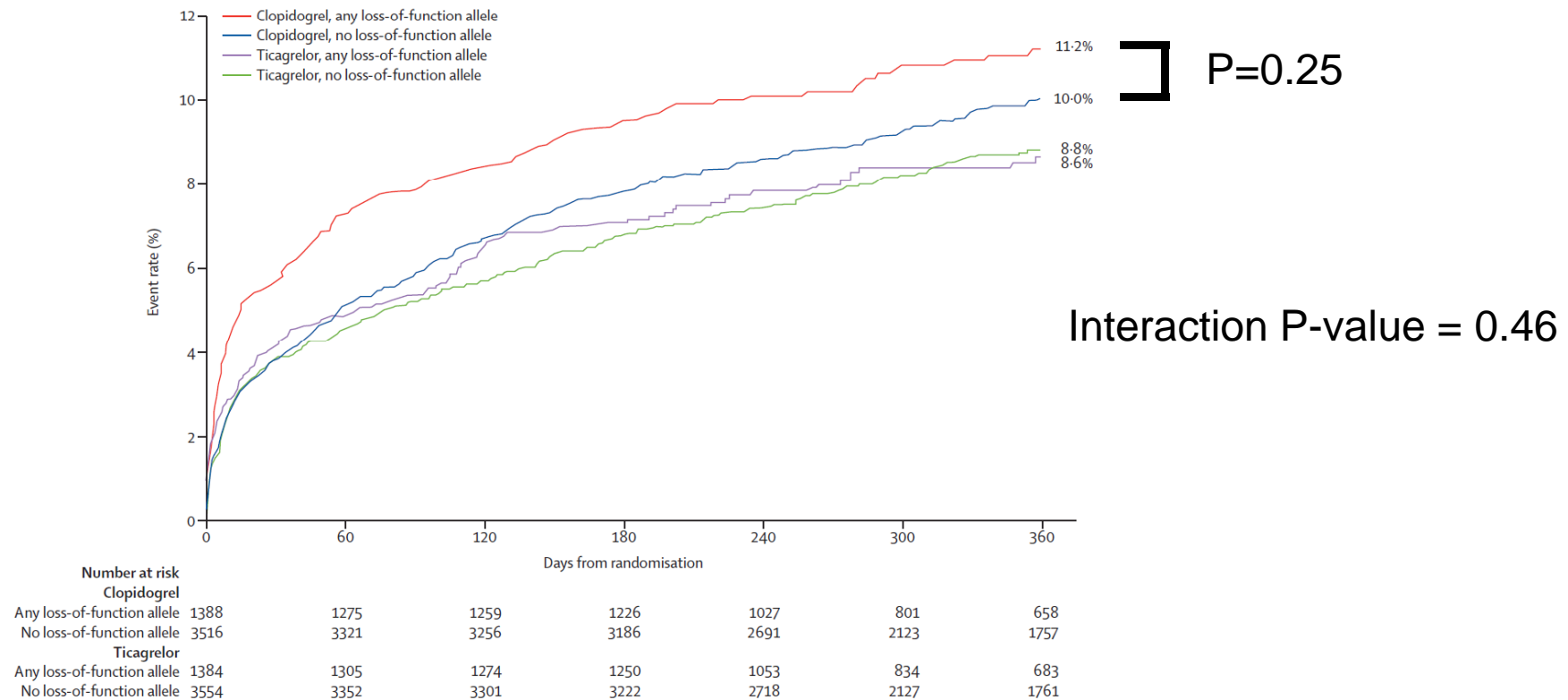
# Does intervention that alters the risk factor lead to clinical benefit?

## PLATO-Genetics Analysis

- 18 624 patients with or without ST-elevation ACS
  - Clopidogrel (75 mg QD) Vs ticagrelor (90 mg BID)
  - Median follow-up of 277 days
- Outcomes
  - First Primary: CV death, MI, Stroke
  - Safety: Major bleed
- 10,285 patients included in the genetic analysis

# Does intervention that alters the risk factor lead to clinical benefit?

## PLATO-Genetics Analysis



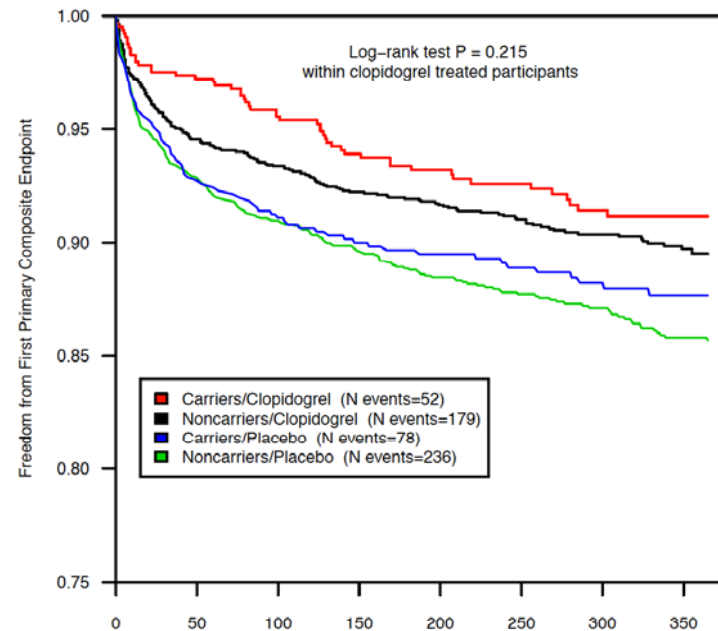
# Does intervention that alters the risk factor lead to clinical benefit?

## CURE-Genetics Analysis

- 12,562 ACS patients without ST-segment elevation
  - Randomized to Clopidogrel (75mg) or Placebo
  - On a background of ASA (75 mg to 325 mg)
  - Average follow-up of 9 month
- Outcomes
  - First Primary: CV death, MI, Stroke
  - Second Primary: First primary, or recurrent ischemia, or UA
  - Safety: Major bleed (life-threatening or not)
- 5,059 patients included in the genetic analysis

# Does intervention that alters the risk factor lead to clinical benefit?

## CURE-Genetics Analysis



| No. at Risk             | Days After Randomization |      |      |      |      |      |     |     |
|-------------------------|--------------------------|------|------|------|------|------|-----|-----|
|                         | 0                        | 50   | 100  | 150  | 200  | 250  | 300 | 350 |
| Carriers Clopidogrel    | 651                      | 632  | 608  | 545  | 484  | 425  | 358 | 297 |
| Noncarriers Clopidogrel | 1886                     | 1778 | 1723 | 1541 | 1352 | 1191 | 960 | 804 |
| Carriers Placebo        | 674                      | 626  | 609  | 551  | 483  | 423  | 356 | 281 |
| Noncarriers Placebo     | 1819                     | 1686 | 1634 | 1456 | 1259 | 1103 | 922 | 774 |

Pare et al. NEJM 2010 363(18): 1704-14



# Other considerations

- Need for further clinical studies
  - Estimates of risk in poor metabolizers versus LOF allele carriers
  - Explain differences in genetic effect between studies
  - Define the best assay and cut-off for PFT
- Need for further genetic studies
  - Heritability ~ 72% but only ~12% explained
- Cost of genetic/PFT testing as compared to alternative treatment
  - Genetic testing can cost up to 500\$

# Conclusion

- The impact of CYP2C19 genotypes/PFT on MACE is real but modest
- There is currently no evidence to support better outcomes from genetically-guided therapy
- Need for randomized trials evaluating genetic testing/PFT as an intervention
  - GRAVITAS, TRIGGER-PCI, ARTIC, etc.
- Future is bright for personalized therapy, but might not (yet) be ready for primetime

# Thank You!

