# Additional findings amongst relatives pursuing oncology multigene panel cascade testing

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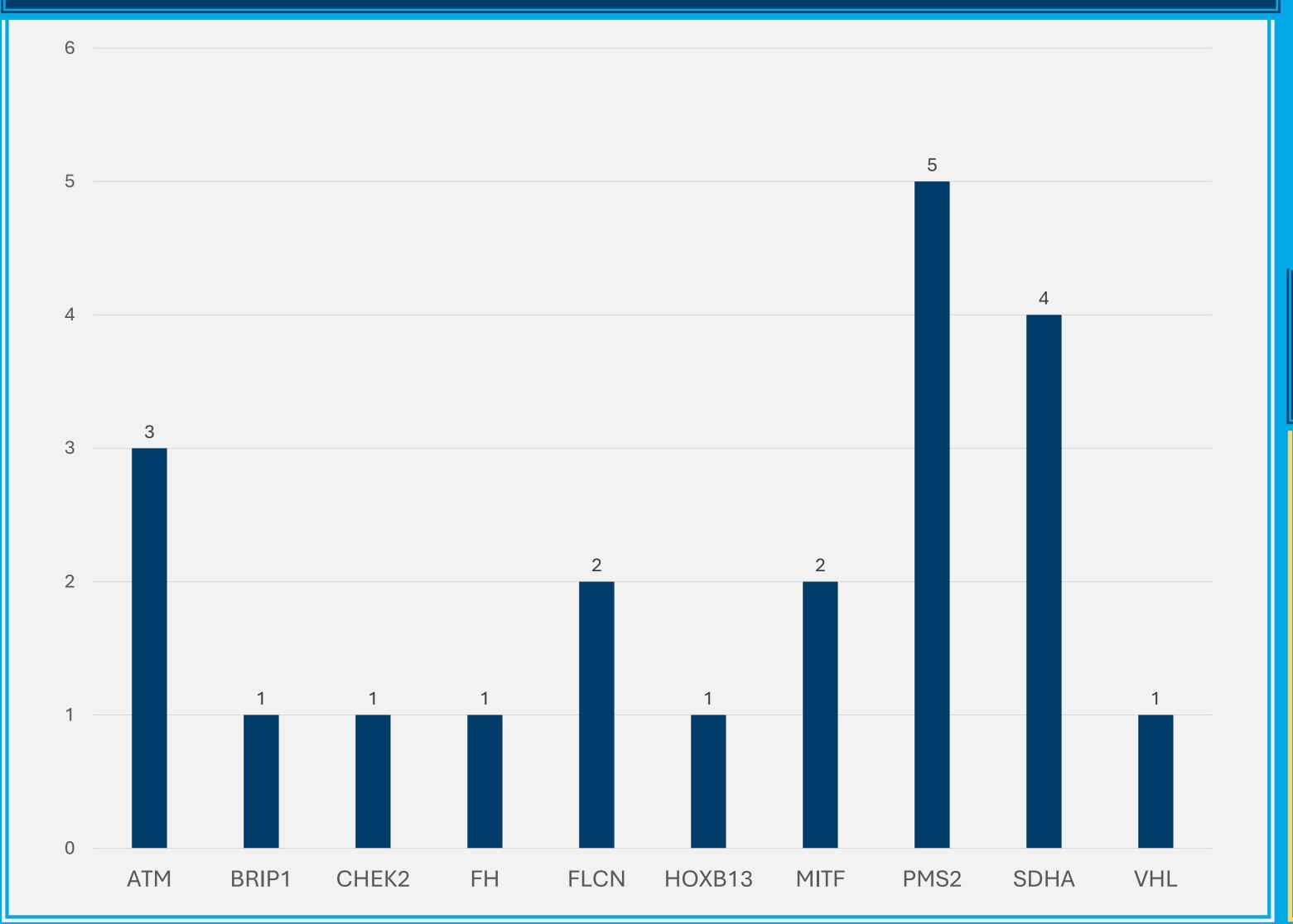
### BACKGROUND & AIMS

- Historically, cascade testing often limited to site specific analysis (SSA)
- Studies investigating outcomes of cascade multigene panel testing (MGPT) as alternative are limited
- Identify additional familial pathogenic variants (AFPVs) and determine if unexpected
- Aid clinicians in determining when cascade MGPT may be more appropriate than SSA

## **METHODS**

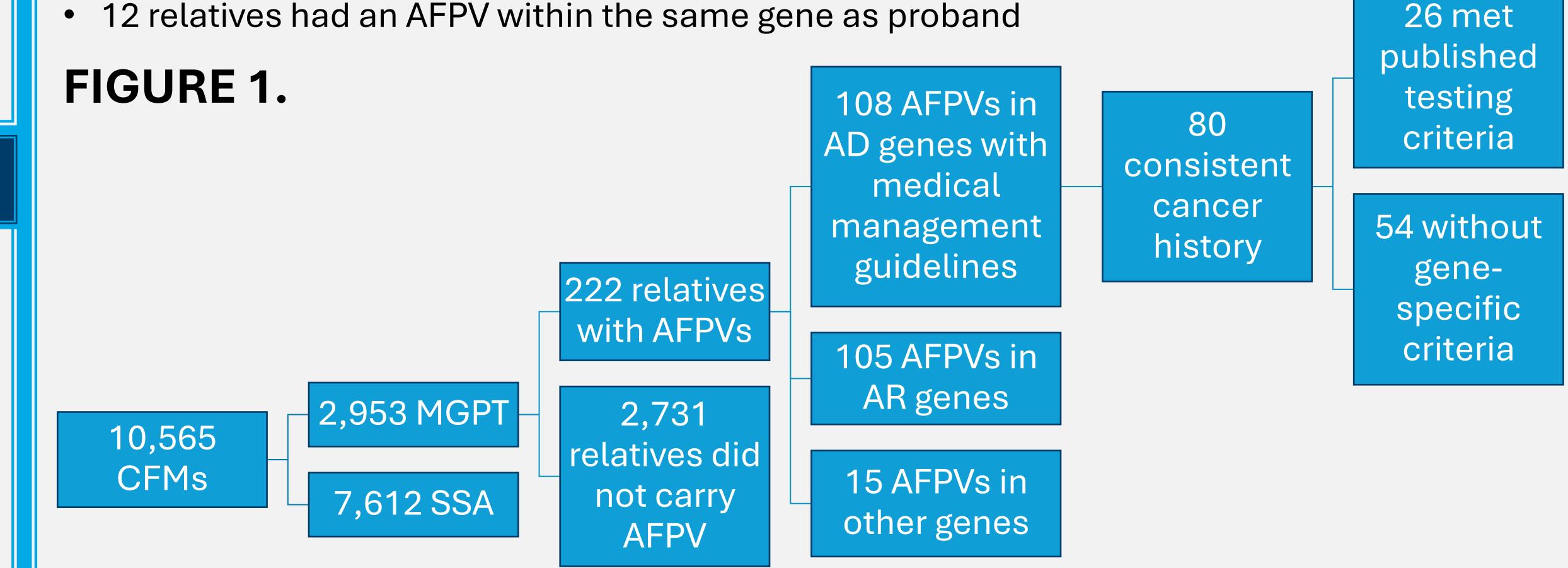
- Retrospective review to identify cascade family members (CFMs) of probands with PVs in BRCA1/2, ATM, CHEK2, PALB2, MLH1, MSH2, MSH6, PMS2, or **EPCAM**
- CFM defined as relative who had testing (SSA, single gene, or MGPT) from 2018-2023, after the proband, that included the proband's impacted gene
- CFM testing of 8+ genes considered MGPT
- Personal and family history provided by ordering clinicians reviewed among individuals with an AFPV to evaluate testing criteria eligibility and management recommendations
- CFM personal and family history was reviewed for published testing criteria eligibility as well as core cancers (e.g. breast, ovarian) to track consistent phenotype in absence of testing criteria

## FIGURE 2. GENES IN WHICH AFPVs IDENTIFIED WITHOUT CONSISTENT HISTORY

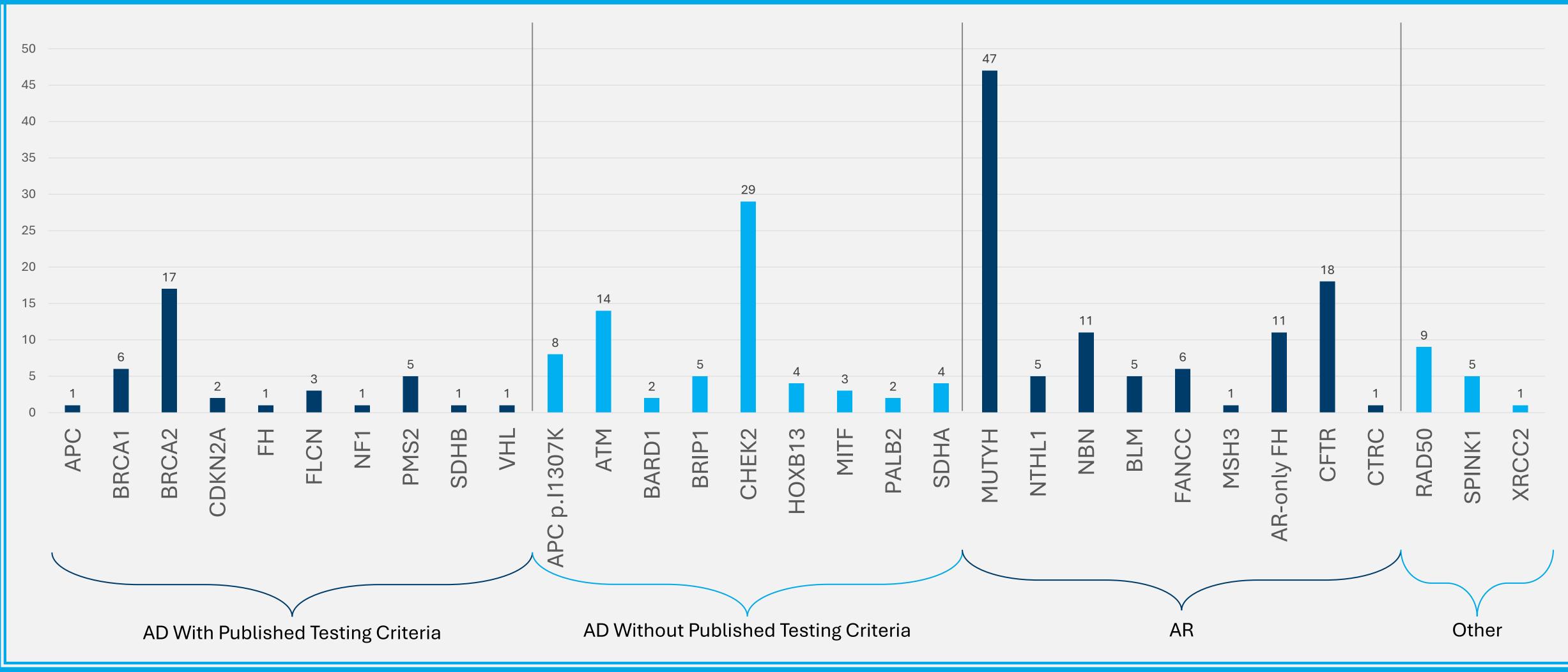


#### RESULTS

- 80 (74.07%) AFPVs consistent with reported personal/family history
  - 26 met published testing criteria (e.g. BRCA1/2)
  - 54 without gene-specific testing criteria (e.g. CHEK2)
- 28 (25.9%) AFPVs impacting medical management identified in individuals without suggestive history (Figure 2)
- 12 relatives had an AFPV within the same gene as proband



### FIGURE 3. AFPVs BY GENE



# TAKE HOME POINTS

- Cascade MGPT increases overall positive rate
- Comprehensive family history collection and pre-test counseling are critical
- 1 in 4 relatives had an AFPV, without suggestive history, impacting their medical management