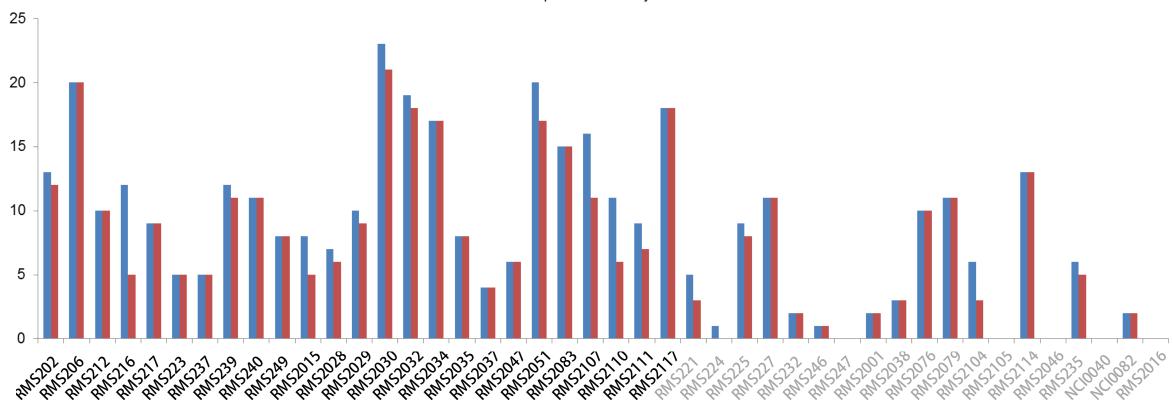
- number of chromosomes/arms with arm-level unexpected copy number and LAF status before normal contamination correction
- number of chromosomes/arms that are corrected to expected status by normal contamination correction



Supplementary Figure S2. Chen et al.