

Supplementary Table 3. Rearrangeable IGHJ genes from Stanford\_S22 dataset as determined from analysis of iHMMune-align, IMGT/V-QUEST+JCTA and IgBLAST alignment results using the method described in (Boyd, et al., 2010).

<b>Rearrangeable IGHJ genes</b>
IGHJ1*01
IGHJ2*01
IGHJ3*02
IGHJ4*02
IGHJ5*02
IGHJ6*02
IGHJ6*03