S2 Fig. Affinity determination for anti-KIR2DL3 mAbs by ELISA. ELISA binding curves were generated using the ECD of KIR2DL3 at 2 µg/mL and serial dilutions of the anti-KIR2DL3 mAbs. Midpoint of the binding curve was used to estimate the affinity ($K_d$) using the Prizm software.