

Figure 1

VK

80.2% identity in 111 residues overlap; Score: 483.0; Gap frequency: 0.0%

Clone47	1	DIVLTQSPASLAVSLGQRATISCRASESDNYGISFMNWFQQKPGQPPKLLIYAASRQGS
HuClone47	1	EIVLTQSPATLSLSPGERATLSCRASESVDNYGISFMNWFQQKPGQAPRLLIYAASRQGS
Clone47	61	GVPARFSGSGGTDFTSLNIHPMEEDDTAMVFCQQSKQVPNTFGGGTKLEIK
HuClone47	61	GIPDRFSGSGGTDFTLTISRLEPEDFAVYCCQQSKQVPNTFGGGTKLEIK

VH

81.5% identity in 119 residues overlap; Score: 530.0; Gap frequency: 0.0%

Clone47	1	QVQLQQPGAEIVRPGASVKLSCKASGYTFSNYLMNWKQRPEDLDWIGRIDPYDGDIDY
HuClone47	1	QVQLVQSGAEVKKPGASVKVSKASGYTFSNYLMNWKQRPEDLDWIGRIDPYDGDIDY
Clone47	61	NQNFKDKAILTVDKSSSTAYMQLSSLTSEDSAVVYCARGYTAYGVDYWGQGTSTVTVSS
HuClone47	61	NQNFKDRVTITRTDSASTAYMELSSLRSEDTAVVYCARGYTAYGVDYWGQGTSTVTVSS

Figure 2

HuCL47, and Clone 47 anti-IL13RA2 IgG to NCI-H1299 cells

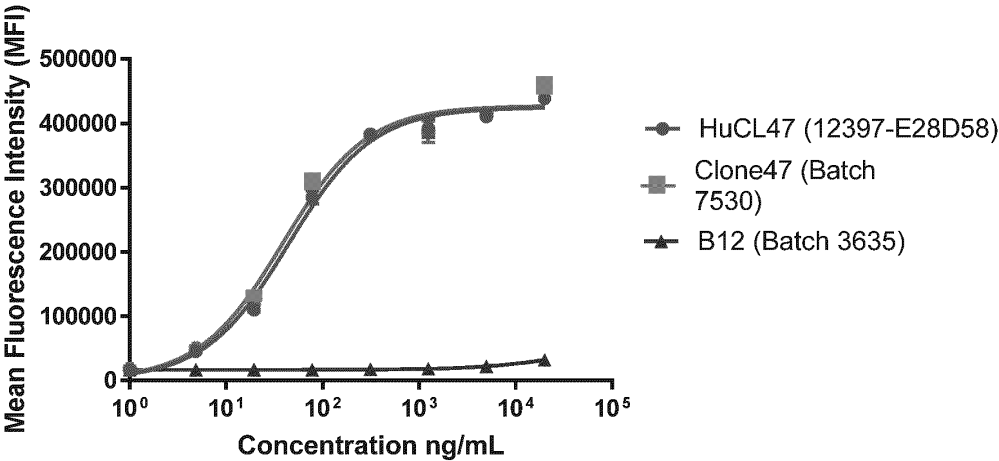


Figure 3

