

[Figure 1]

SEQ ID NO: 1 - Amino acid sequence of a heavy chain of the anti-TROP2 antibody

MKHLWFFLLLVAAPRWLSQVQLVQSGAEVKKPGASVKVSCKASGYTFTTAGMQWRQAPGQ
GLEWMGWINTHSGVPKYAEDFKGRVTISADTSTSTAYLQLSSLKSEDTAVYYCARSFGSSY
WYFDVWGQGTLVTVSSASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGAL
TSGVHTFPAVLQSSGLYSLSSVTVPSSSLGTQTYICNVNHKPSNTKVDKRVEPKSCDKTHT
CPPCPAPELLGGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNA
KTKPREEQYNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVY
TLPPSREEMTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTPPVLDSDGSFFLYSKLT
VDKSRWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK

Signal sequence (1-19), Variable region (20-140),
Constant region (141-470)

[Figure 2]

SEQ ID NO: 2 - Amino acid sequence of a light chain of the anti-TROP2 antibody

MVLQTQVFISLLLWISGAYGDIQMTQSPSSLSASVGDRVTITCKASQDVSTAVAWYQQKPGK
APKLLIYSASYRYTGVPSTRFSGSGSGTDFTLTISLQPEDFAVYYCQQHYITPLTFGQGTKL
EIKRTVAAPSVFIFPPSDEQLKSGTASVVCLLNNFYPREAKVQWKVDNALQSGNSQESVTEQ
DSKDSTYLSSTLTLSKADYEKHKVYACEVTHQGLSSPVTKSFNRGEC

Signal sequence (1-20), Variable region (21-129),
Constant region (130-234)

[Figure 3]

SEQ ID NO: 3 - Amino acid sequence of a heavy chain of the anti-B7-H3 antibody

MKHLWFFLLLVAAPRWLSQVQLVQSGAEVKKPGSSVKVSCASGYTFTNYVMHWVRQAPGQ
GLEWMGYINPYNDVKYNEKFGRVTITADESTSTAYMELSSLRSEDNAVYYCARWGYYGSP
LYYFDYWGGTLTVSSASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGA
LTSGVHTFPAVLQSSGLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKRVEPKSCDKTH
TCPPCPAPELLGGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHN
AKTKPREEQYNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQV
YTLPPSREEMTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKL
TVDKSRWQQGNVFCFSVMHEALHNHYTQKSLSLSPGK

Signal sequence (1-19), Variable region (20-141),
Constant region (142-471)

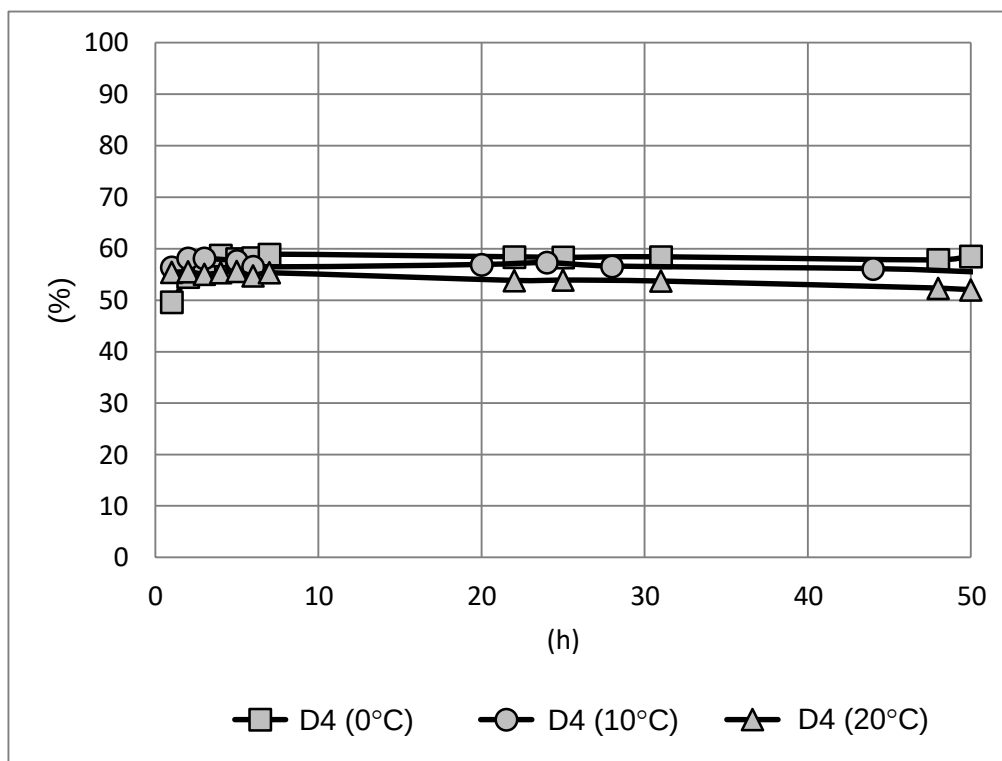
[Figure 4]

SEQ ID NO: 4 - Amino acid sequence of a light chain of the anti-B7-H3 antibody

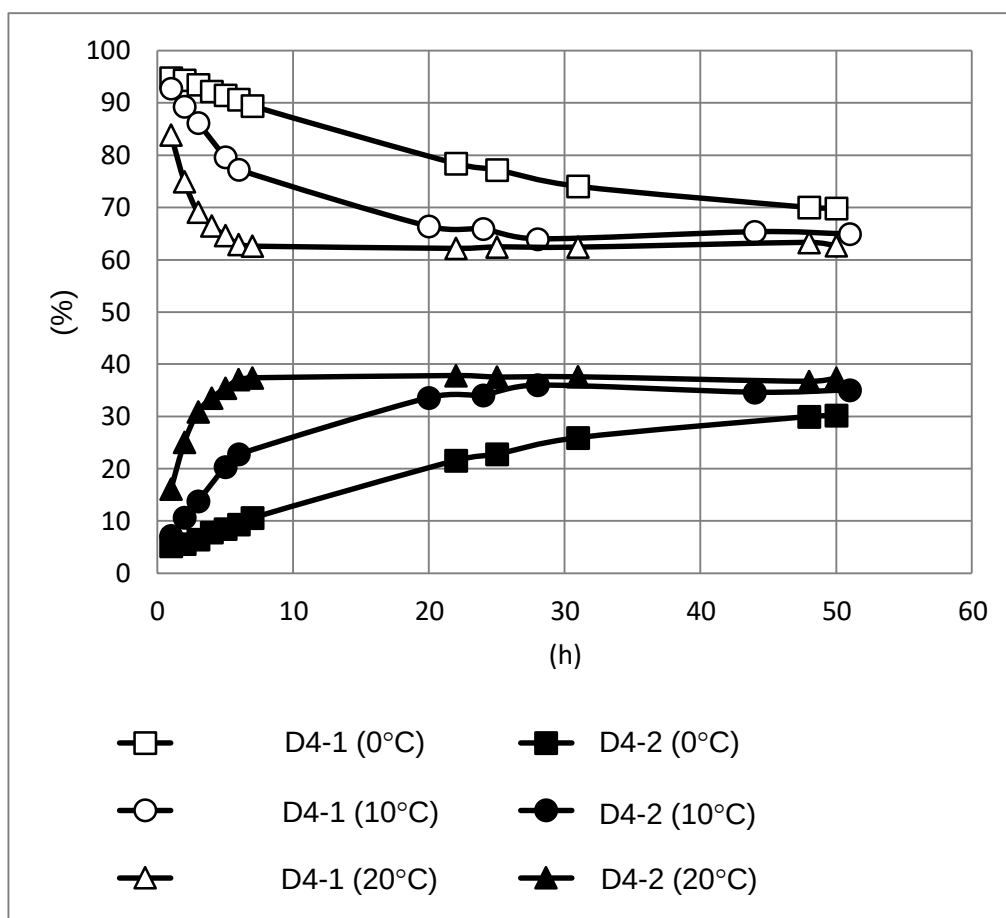
MVLQTQVFISLLLWISGAYGEIVLTQSPATLSLSPGERATLSCRASSRLIYMHWYQQKPGQA
PRPLIYATSNLASGIPARFSGSGSGTDFTLTISSLEPEDFAVYYCQQWNSNPPTFGQGTKVE
IKRTVAAPSVFIFPPSDEQLKSGTASVVCLLNNFYPREAKVQWKVDNALQSGNSQESVTEQD
SKDSTYLSSTLTLSKADYEKHKVYACEVTHQGLSSPVTKSFNRGEC

Signal sequence (1-20), Variable region (21-128),
Constant region (129-233)

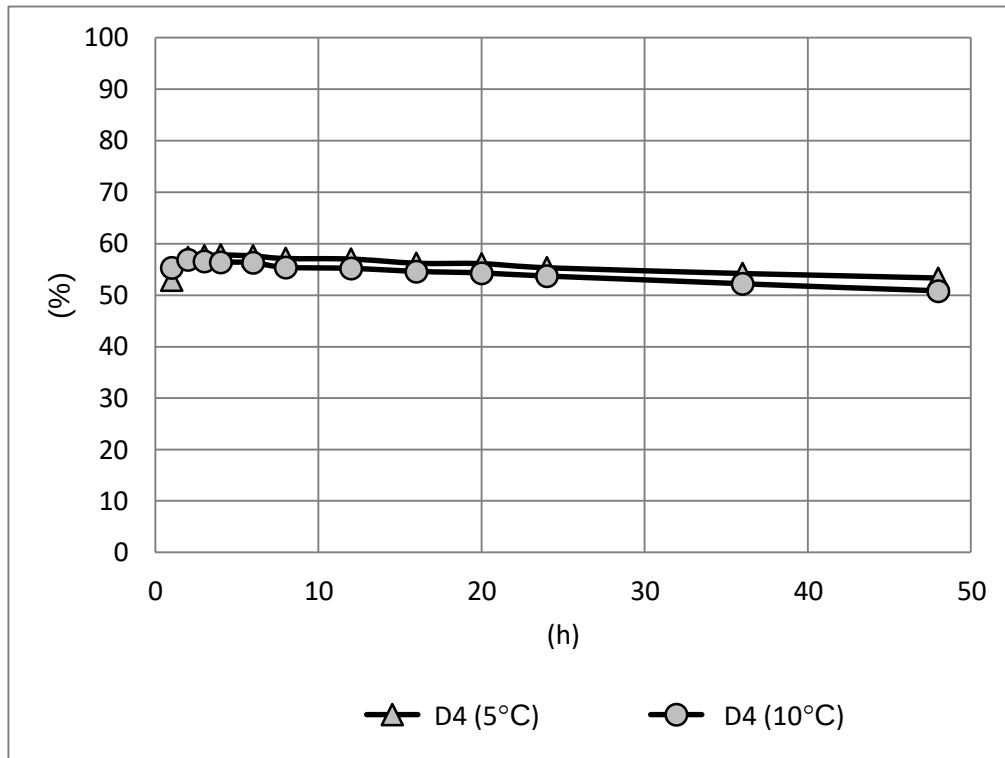
[Figure 5]



[Figure 6]



[Figure 7]



[Figure 8]

