

In the article "High-frequency representation of a single V_H gene in the expressed human B cell repertoire" by A. K. Stewart, C. Huang, B. D. Stollar, and R. S. Schwartz (February 1993, 177:409), Figure 5 was printed incorrectly. The correct version of Figure 5 and its legend appear below.

CLONE	N	D	N	JH	
AL1.1		GATATTGTAcTGGTGGT gta TGC	GGGGGATGAT	GCTTTTGATATC	TGGGG
AL1.2	GATATCG	CCAGCTGCTAT	AAAAGT	TACTgCGcTATGGACGTC	TGGGG
AL1.3	GATAAGGC	GTATTATGA	TACTCCA	TTGACTAC	TGGGG
TK1	GATCCGGTCCA	CAGCTGG	GCC	AACTGGTTCGACcCC	TGGGG
AL3.1	GGATCGGGCGAAAACACCAAAATGACTACGGTG	ACTACTTTC CGACATAC		TACTTTGACTAC	TGGGG
AL3.2	CTATCCGCCG	ATAGTGGCTACGATT	CGGGGGAA	TACTTTGACTAC	TGGGG
AL3.4	GATCCGTCCGCT	TACAGTAAcT	CAGTGGGTAGA	TTGACTAC	TGGGG
AL3.7	CGGATCCCGC	CAGCAGacGG ACGATA C		TACTTTGACTAC	TGGGG
AL3.10	GATACCCGAGAGGAA	GATTTTGGAGTGGTTATTATA		ACTACTACGGTATGGACGTC	TGGGG
AL3.11	GGCCCG	TATTTGACTGGTTA	AGG		
AL3.16	TGTAGCGAACTC	CGATATTTGACTGGTcAgT		TGACTAC	TGGGG

Figure 5. Nucleotide sequences of the CDR3 segments of the 12 V_H18/2 hybridizing plaques. Areas with identity to known germline gene segments are shown in bold type, mutations in plain type. These sequence data are available from EMBL/GenBank/DBJ under accession numbers X67060–X67074.