

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		DLE2 GATATTGTAcTGGTGGTgtaTGC	GGGGGATGAT	GCTTTTGATATC	TGGGG
AL1.2	GATATCG	DLE4 CCAGCTGCTAT	AAAAGT	TACTgCGcTATGGACGTC	TGGGG
AL1.3	GATAAGGC	DXP2 (rev) GTATTATGA	TACTCCAA	TTGACTAC	TGGGG
TK1	GATCCGGTCCA	DM1 CAGCTGG	GCC	AACTGGTTCGACcCC	TGGGG
AL3.1	GGATCGGGCGAAAACACCAAATGACTACGGTG	DLE5 ACTACTTTC	CGACATAC	TACTTTGACTAC	TGGGG
AL3.2	CTATCCGCCG	DK1 ATAGTGGCTACGATT	CGGGGGAA	TACTTTGACTAC	TGGGG
AL3.4	GATCCGTCGCCT	DA1 TACAGTAAcT	CAGTGGGTAGA	TTGACTAC	TGGGG
AL3.7	CGGATCCCGC	DM1 CAGCAGacGG	DXP1 ACGATA C	TACTTTGACTAC	TGGGG
AL3.10	GATACCCGAGAGGAA	DXP4 GATTTTGGAGTGGTTATTATA		ACTACTACGGTATGGACGTC	TGGGG
AL3.11	GGCCCG	DXP1 TATTTTGACTGGTTA	AGG		
AL3.16	TGTAGCGAACTC	DXP1 CGATATTTTGACTGGTcAgT		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.