

Supplemental list of sequence logos

This file presents KLID-base sequence logos of complete mixtures generated throughout the study. Shown are KLID-based sequence logos for two to eight-component mixtures (M02 to M08). The file is divided into two parts:

Part 1: mixture component logo for virus-representative (pages 2 - 10)

Component sequence logos of representative IS mixtures for HIV, HTLV, MLV, MVV, PFV, PFV, and ASLV. IS sets used for the mixture estimations are named in following table:

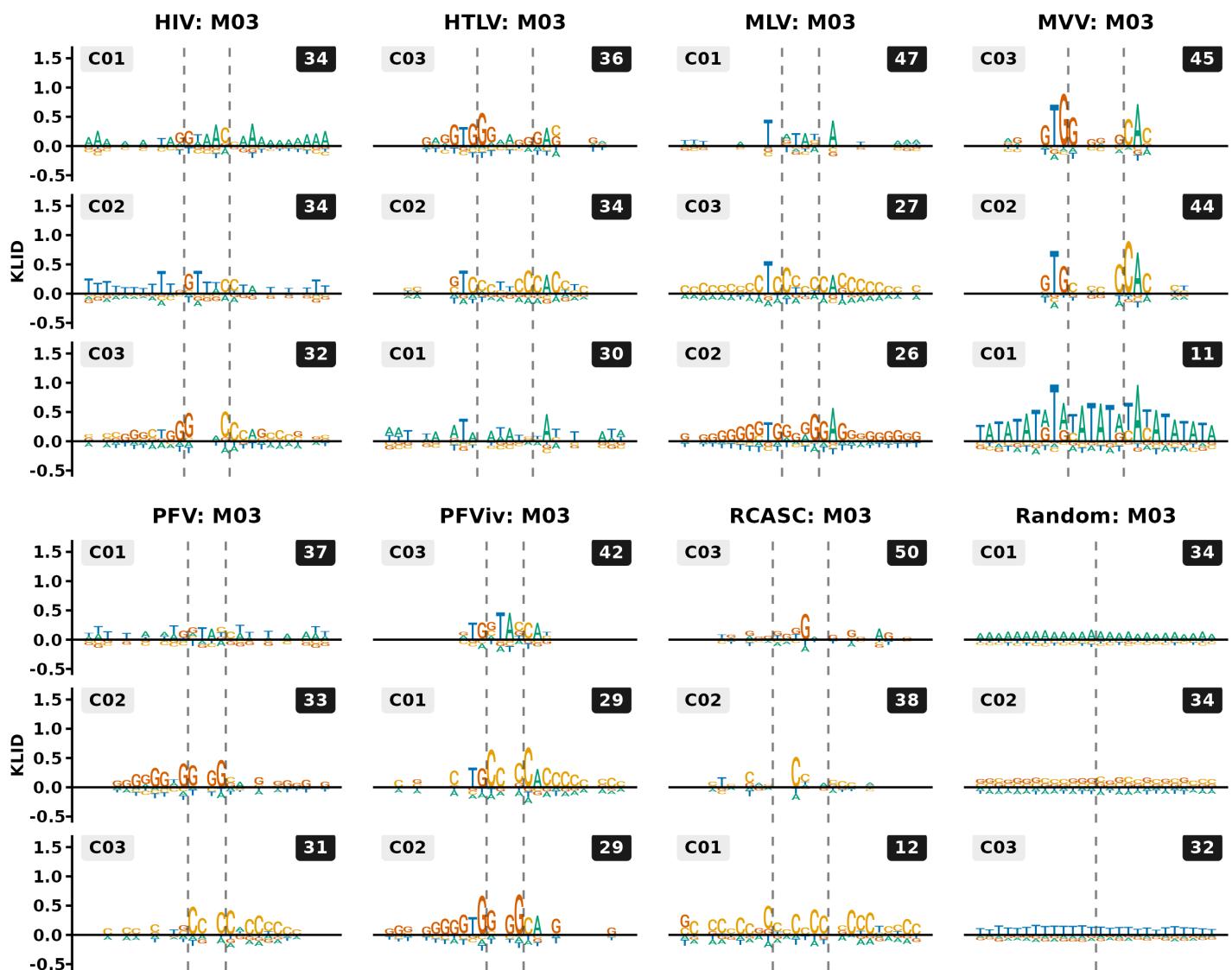
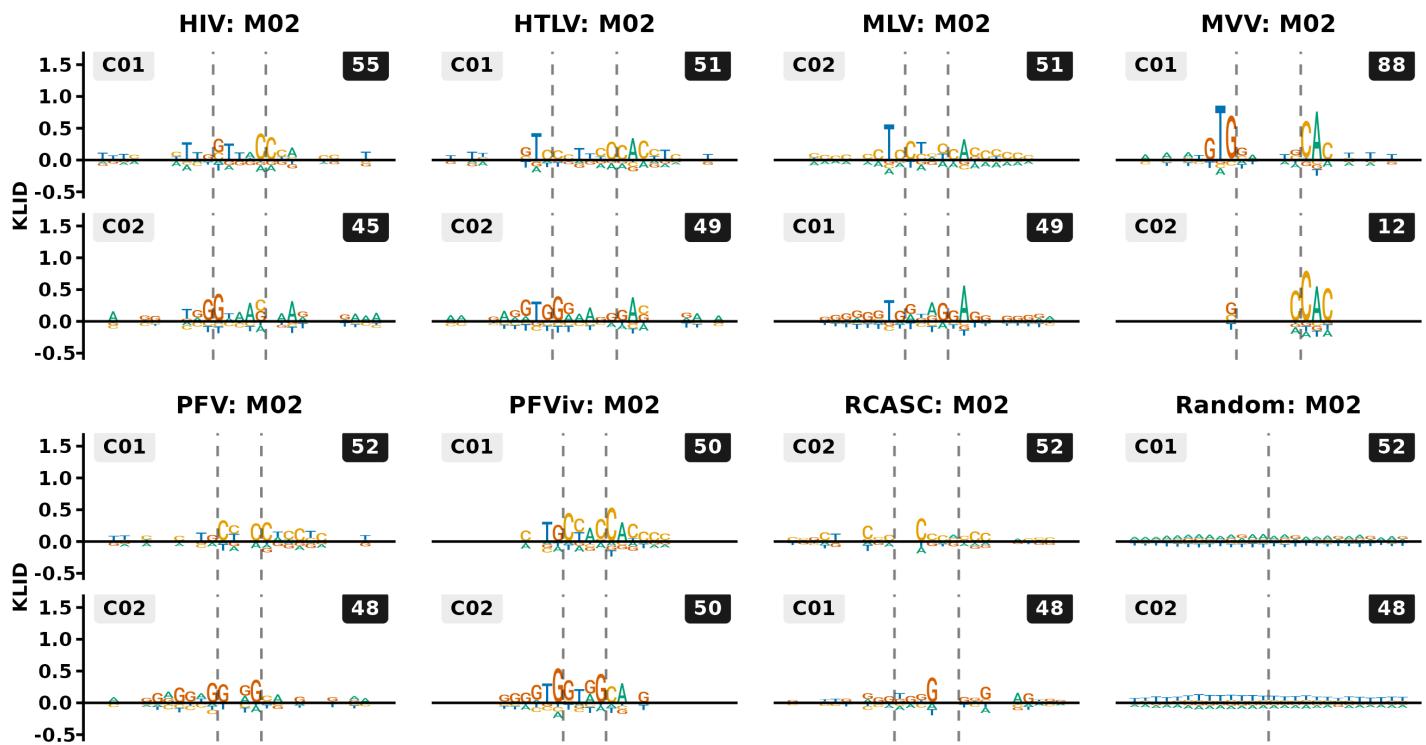
Virus	Cells	#IS	Publication
HTLV	Jurkat	4521	Kirk et al. 2016
HIV	Jurkat	161470	Zhyvoloup et al. 2017
MLV	Human CD34+	57991	De Ravin et al. 2014
MVV	HEK293T	411721	Ballandras-Colas et al 2022
PFV	HT1080	55384	Lesbats et al. 2017
ASLV	CEF	2096	Malhotra et al. 2017

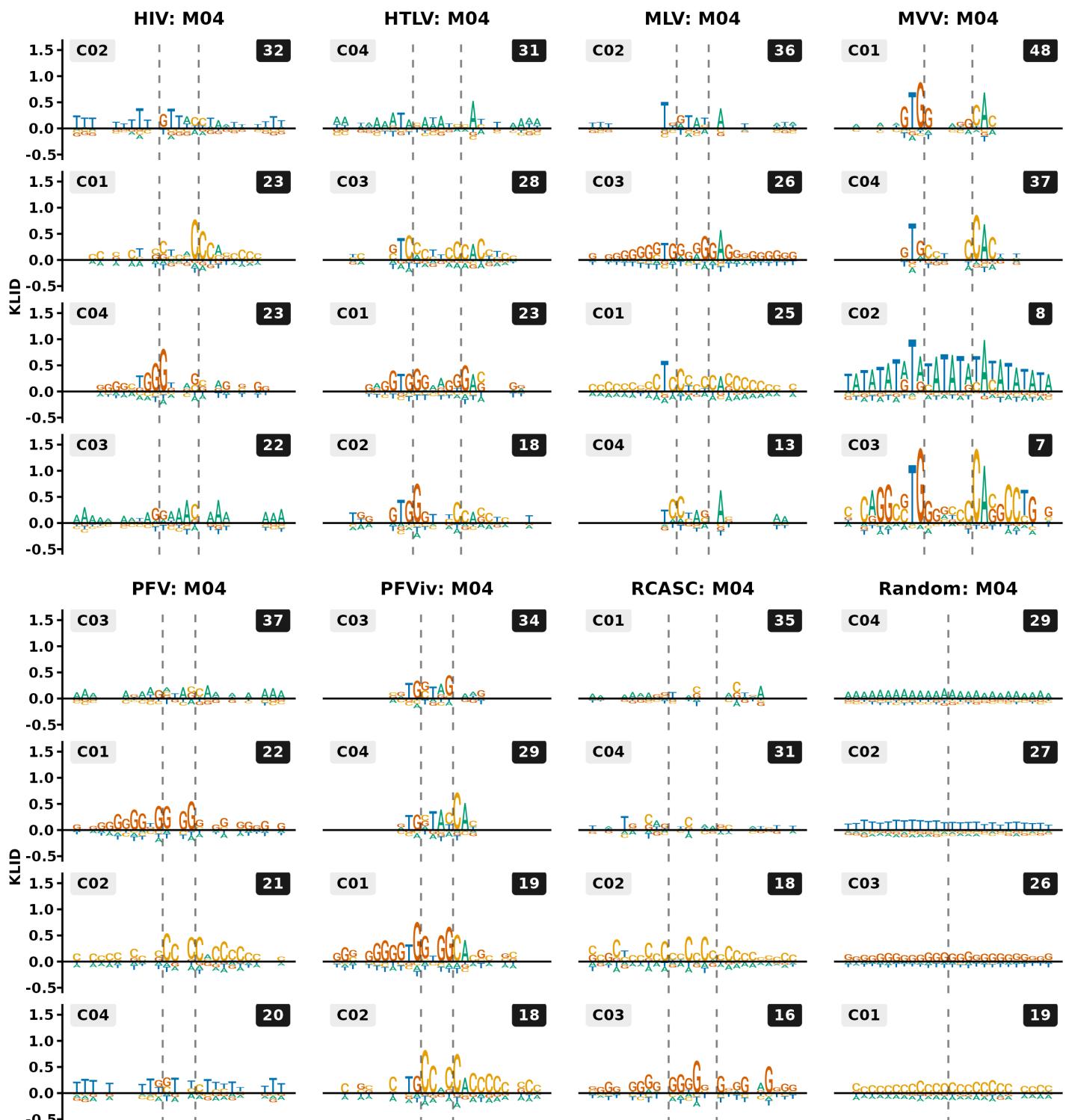
Part 2: Comparing mixture components for HIV and HTLV IS data sets (pages 11 - 19)

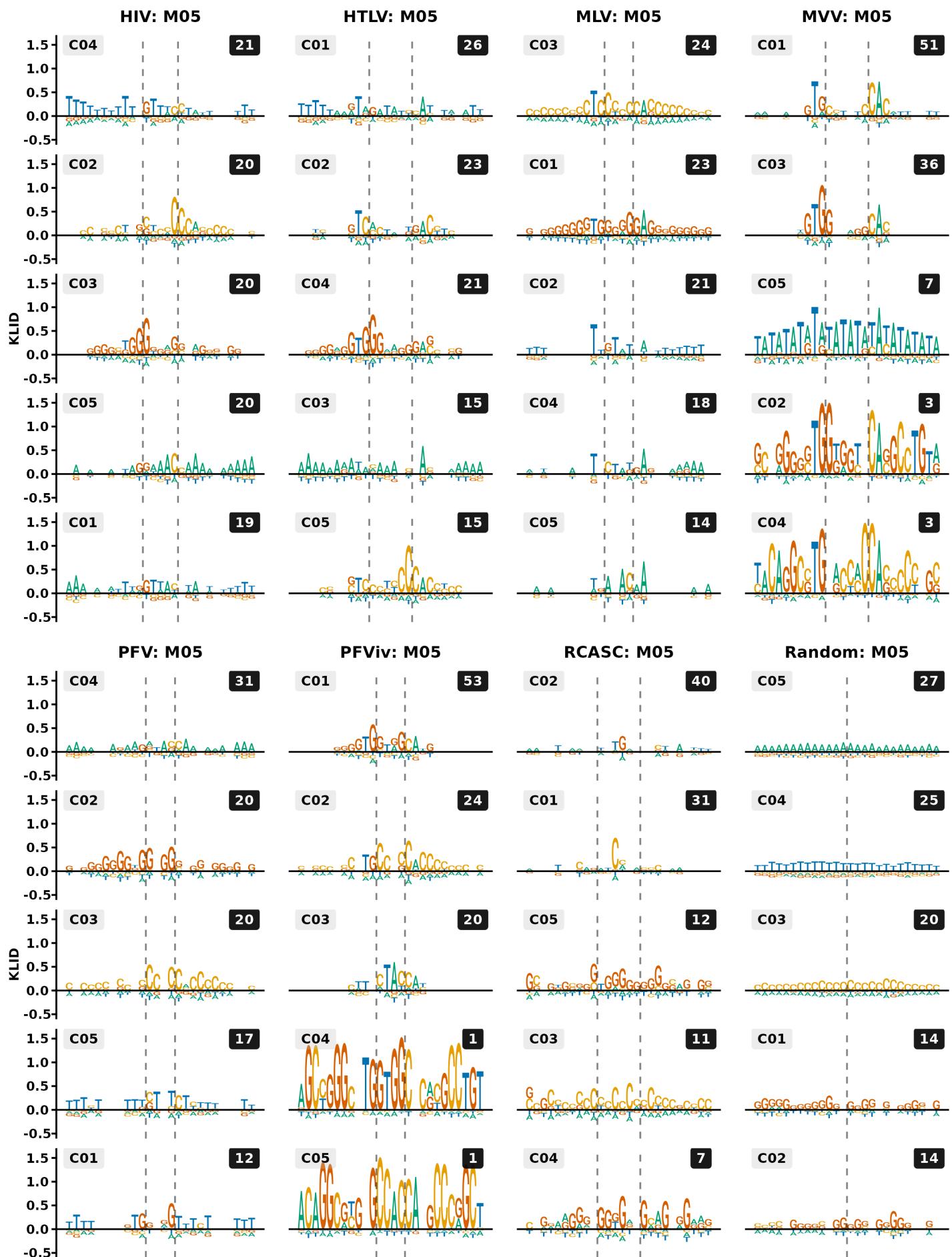
Side-by-side comparison of estimated mixtures of experimentally independent IS sets representing similar viruses. Data sets used are named in the following table:

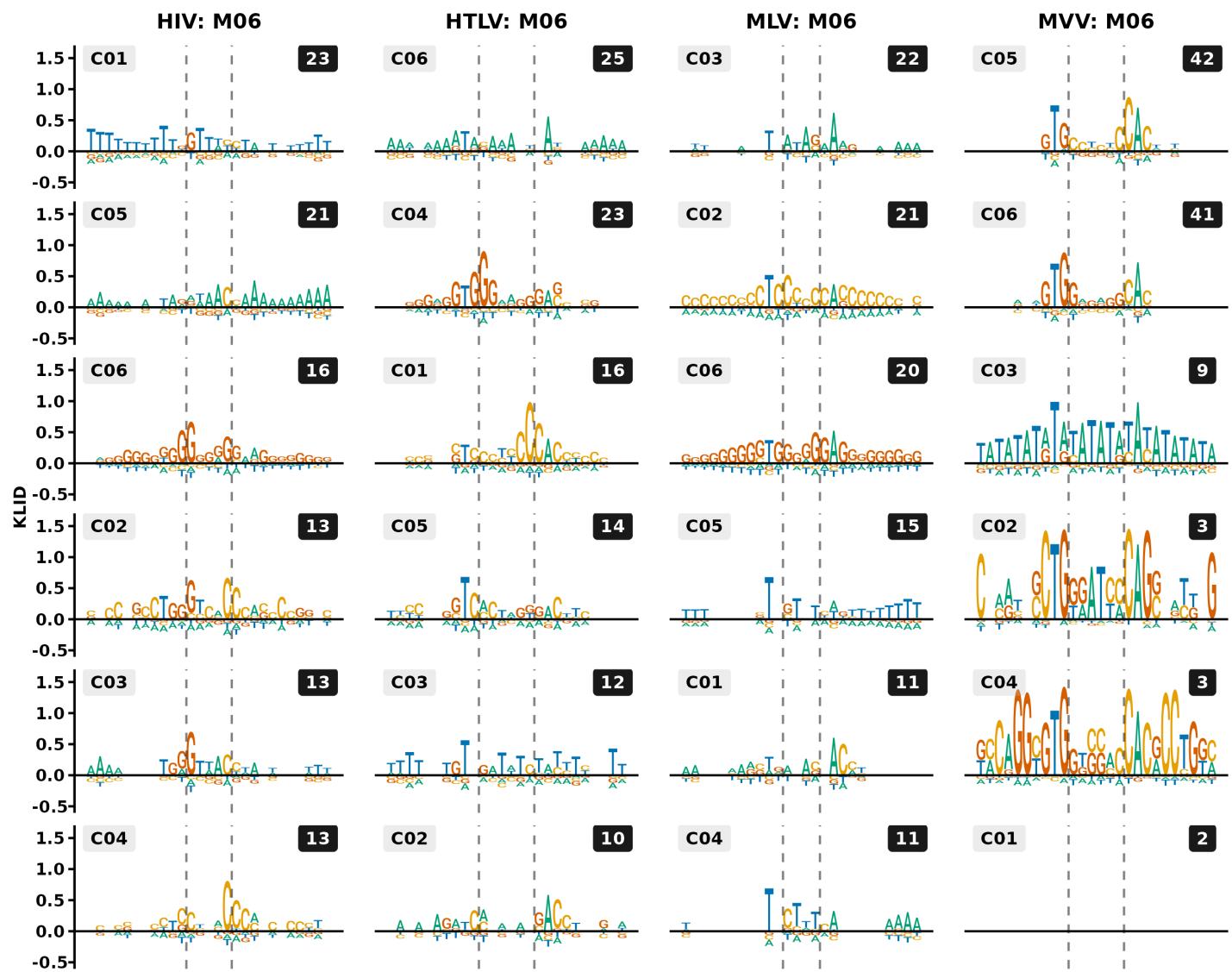
Virus	Cells	#IS	Publication
HTLV	Jurkat	4521	Kirk et al. 2016
HTLV	patient PBMC	162760	Melamed et al. 2022
HTLV	cell culture (various)	235150	Melamed et al. 2022
HIV	Jurkat	161470	Zhyvoloup et al. 2017
HIV	SupT1	16803	Vasant et al. 2020
HIV	patient PBMC	22225	Melamed et al. 2022
HIV	cell culture (various)	33081	Melamed et al. 2022

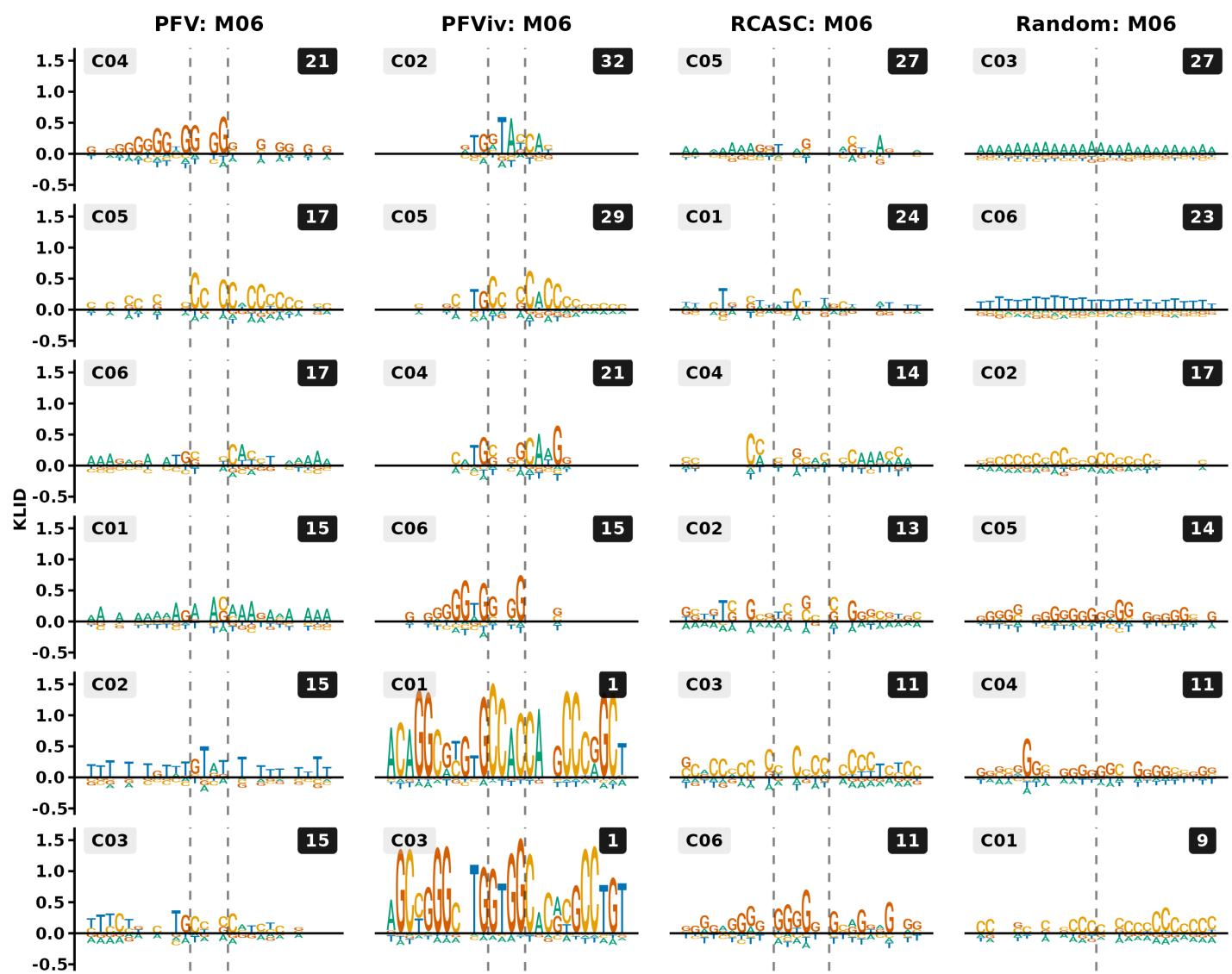
Part 1

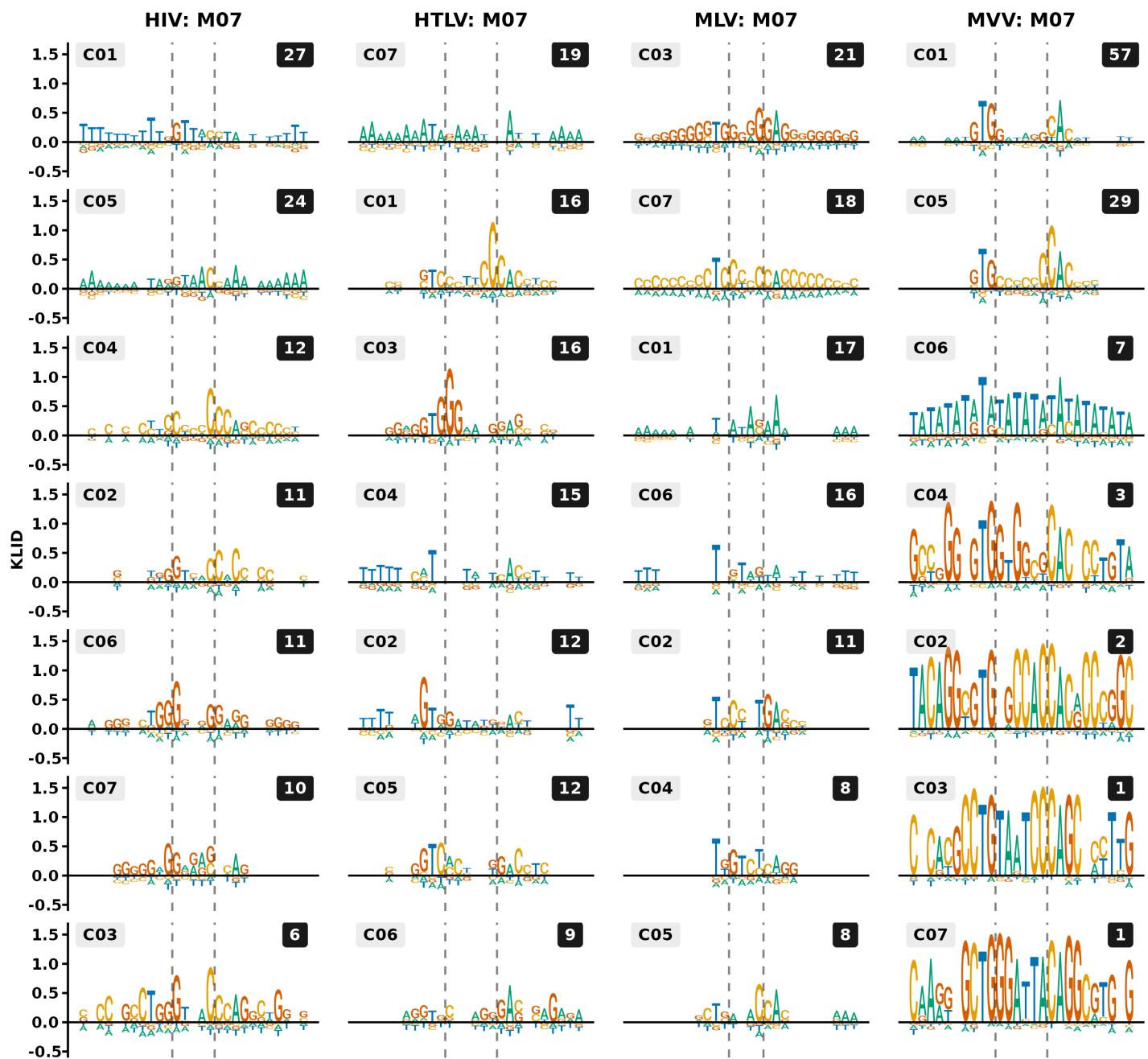


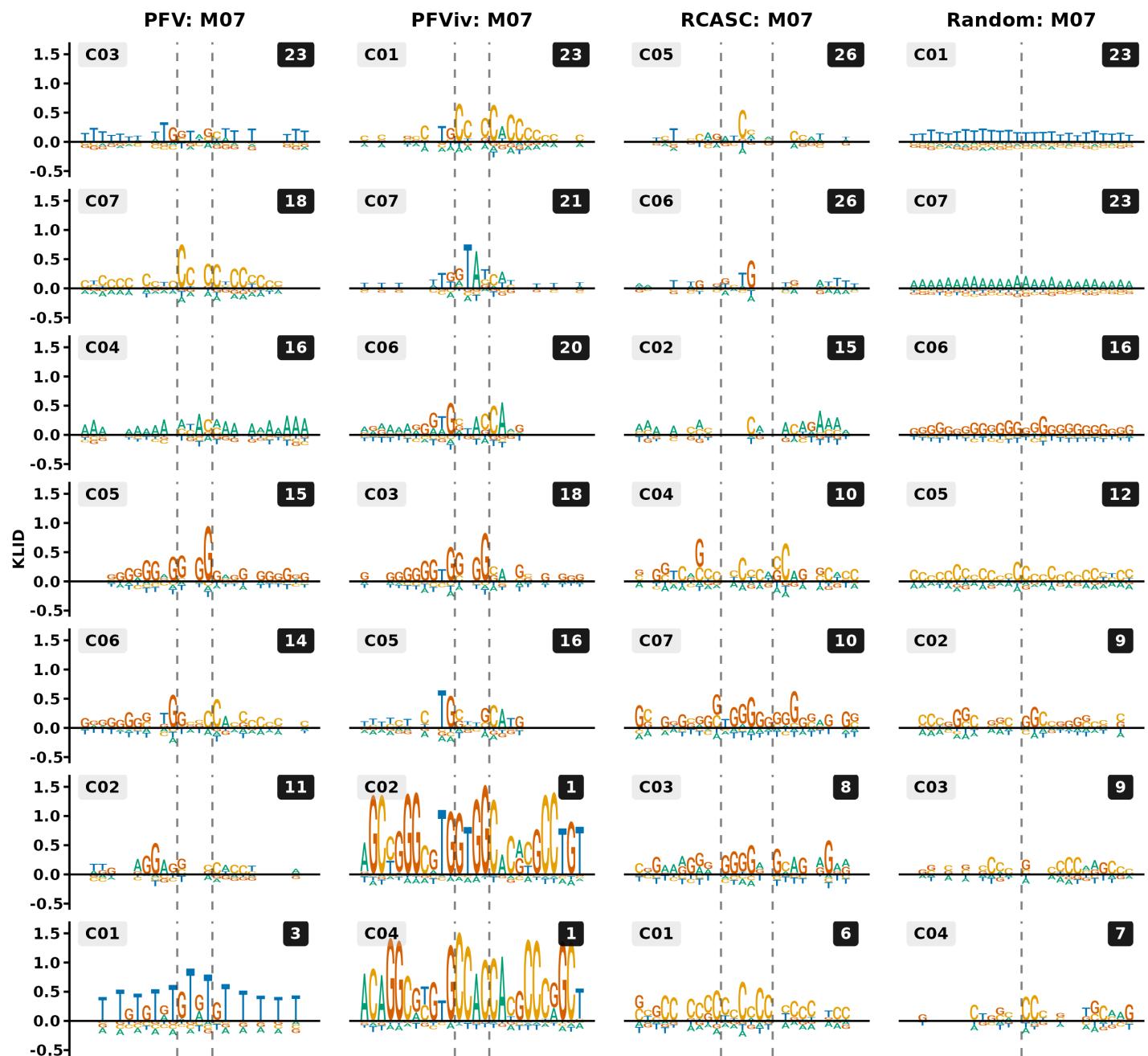


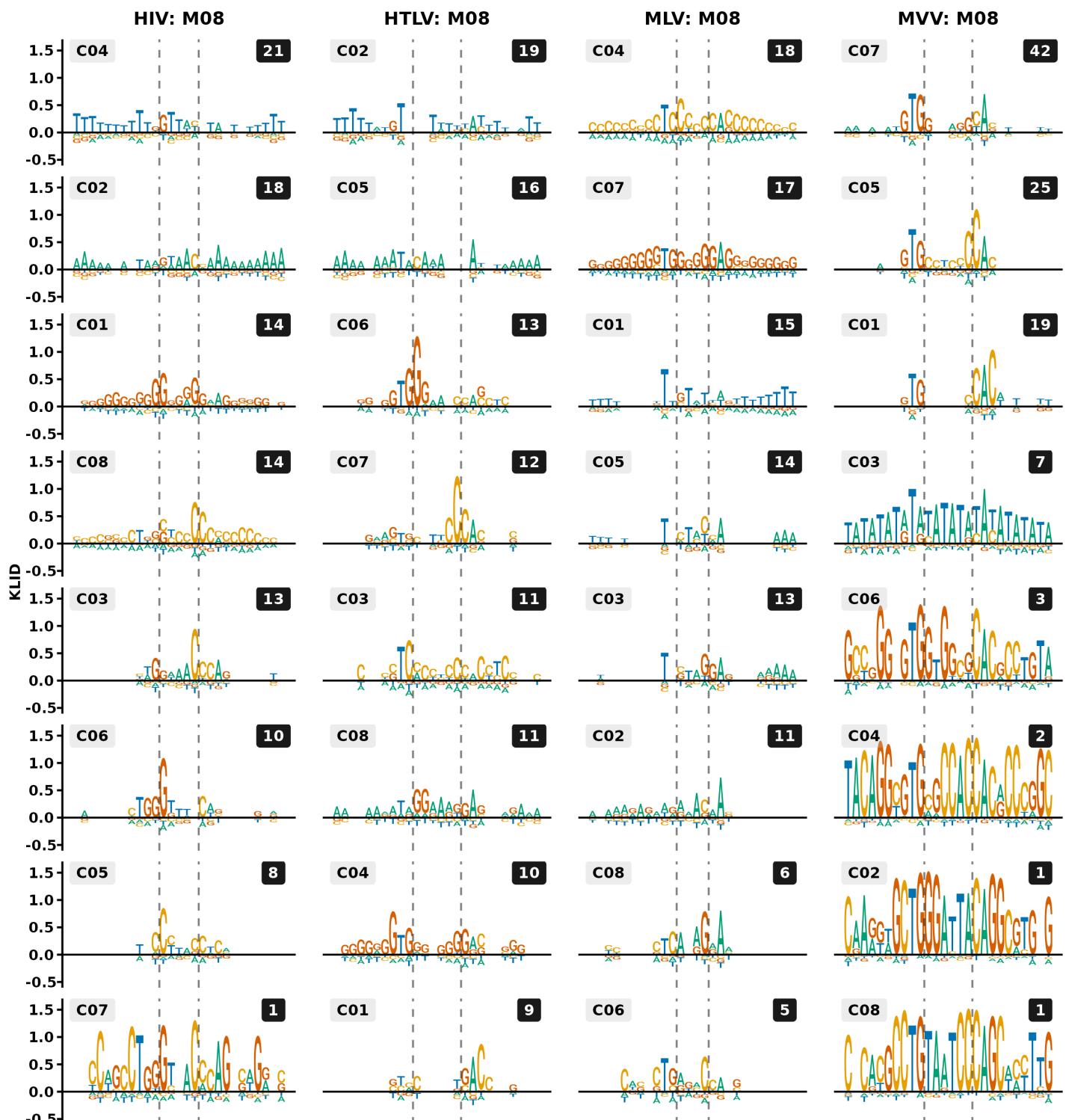


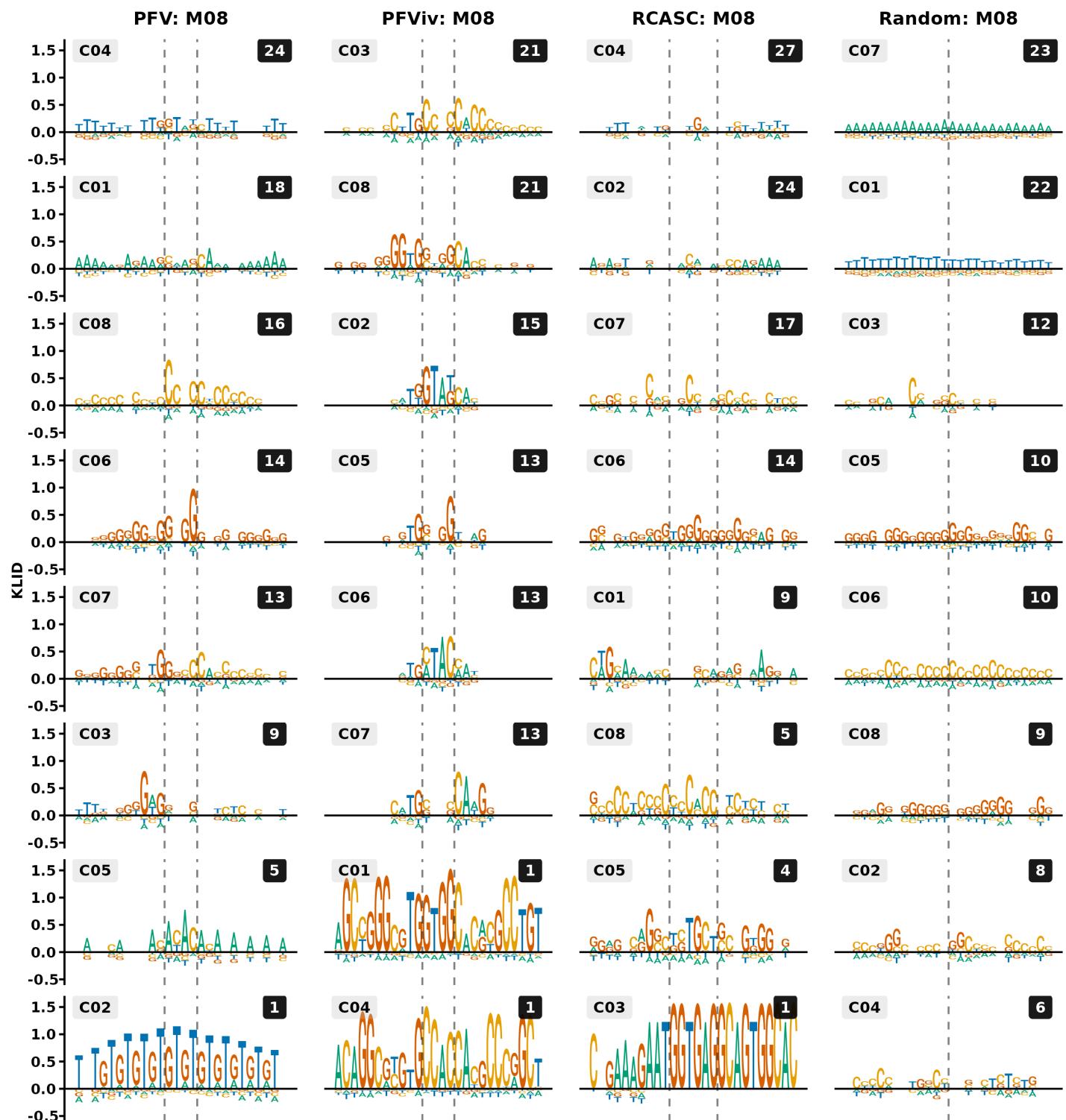












Part 2

