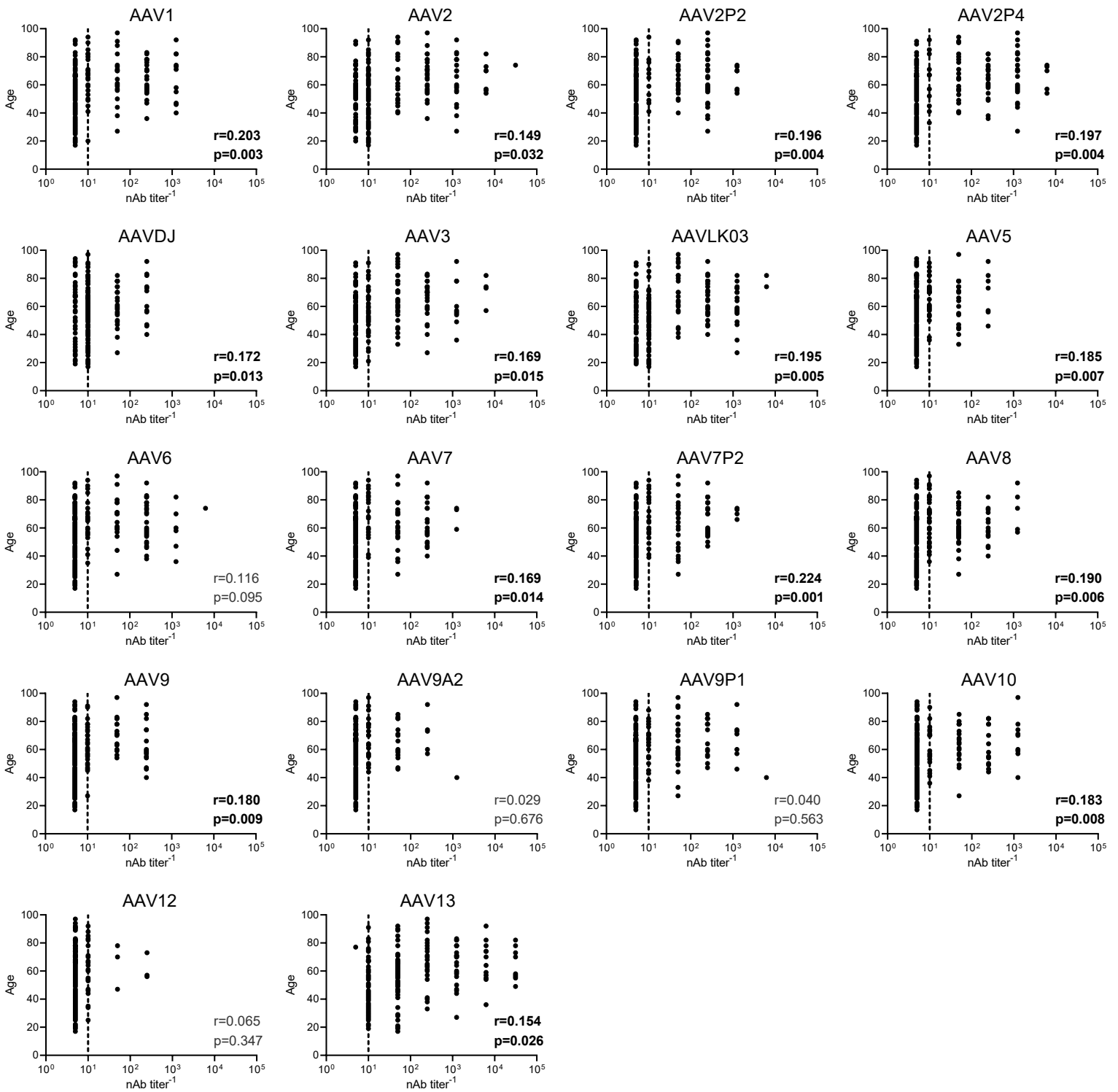


# Supplementary Figure S1

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AAV12/1-743	1	MAADGYLPLDWLEDNLSEGIREMWWALKPGFAPKPKANQQHDDARGLVLPQNYLQFNGLDRGPEVNRADAAAALHDKAYDQDLKAGDNPYLYKNHADAEFQERL	104
AAV9/1-737	1	MAADGYLPLDWLEDNLSEGIREMWWALKPGFAPKPKANQQHDDARGLVLPQNYLQFNGLDRGPEVNRADAAAALHDKAYDQDLKAGDNPYLYKNHADAEFQERL	104
AAV1/1-737	1	MAADGYLPLDWLEDNLSEGIREMWWALKPGFAPKPKANQQHDDARGLVLPQNYLQFNGLDRGPEVNRADAAAALHDKAYDQDLKAGDNPYLYKNHADAEFQERL	104
AAV6/1-737	1	MAADGYLPLDWLEDNLSEGIREMWWALKPGFAPKPKANQQHDDARGLVLPQNYLQFNGLDRGPEVNRADAAAALHDKAYDQDLKAGDNPYLYKNHADAEFQERL	104
AAV7/1-738	1	MAADGYLPLDWLEDNLSEGIREMWWALKPGFAPKPKANQQHDDARGLVLPQNYLQFNGLDRGPEVNRADAAAALHDKAYDQDLKAGDNPYLYKNHADAEFQERL	104
AAV13/1-734	1	MTDGYLPLDWLEDNLSEGIREMWWALKPGFAPKPKANQQHDDARGLVLPQNYLQFNGLDRGPEVNRADAAAALHDKAYDQDLKAGDNPYLYKNHADAEFQERL	103
AAV36/1-737	1	MAADGYLPLDWLEDNLSEGIREMWWALKPGFAPKPKANQQHDDARGLVLPQNYLQFNGLDRGPEVNRADAAAALHDKAYDQDLKAGDNPYLYKNHADAEFQERL	104
AAV/LK03/1-737	1	MAADGYLPLDWLEDNLSEGIREMWWALKPGFAPKPKANQQHDDARGLVLPQNYLQFNGLDRGPEVNRADAAAALHDKAYDQDLKAGDNPYLYKNHADAEFQERL	104
AAV8/1-739	1	MAADGYLPLDWLEDNLSEGIREMWWALKPGFAPKPKANQQHDDARGLVLPQNYLQFNGLDRGPEVNRADAAAALHDKAYDQDLKAGDNPYLYKNHADAEFQERL	104
AAV10/1-739	1	MAADGYLPLDWLEDNLSEGIREMWWALKPGFAPKPKANQQHDDARGLVLPQNYLQFNGLDRGPEVNRADAAAALHDKAYDQDLKAGDNPYLYKNHADAEFQERL	104
AAV2/1-736	1	MAADGYLPLDWLEDNLSEGIREMWWALKPGFAPKPKANQQHDDARGLVLPQNYLQFNGLDRGPEVNRADAAAALHDKAYDQDLKAGDNPYLYKNHADAEFQERL	104
AAV/DJ/1-738	1	MAADGYLPLDWLEDNLSEGIREMWWALKPGFAPKPKANQQHDDARGLVLPQNYLQFNGLDRGPEVNRADAAAALHDKAYDQDLKAGDNPYLYKNHADAEFQERL	104
AAV5/1-725	104	ADTDSFGGNGLRVFAQAKRRLLEPLGLVVEAAKTAAPKRRPVEQSPQEPDSSASISGKTSSTSSDAEAGPSGSLQLQIAQPASSL	188
AAV12/1-743	105	ADTDSFGGNGLRVFAQAKRRLLEPLGLVVEAAKTAAPKRRPVEQSPQEPDSSASISGKTSSTSSDAEAGPSGSLQLQIAQPASSL	201
AAV9/1-737	105	ADTDSFGGNGLRVFAQAKRRLLEPLGLVVEAAKTAAPKRRPVEQSPQEPDSSASISGKTSSTSSDAEAGPSGSLQLQIAQPASSL	198
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AAV6/1-737	105	ADTDSFGGNGLRVFAQAKRRLLEPLGLVVEAAKTAAPKRRPVEQSPQEPDSSASISGKTSSTSSDAEAGPSGSLQLQIAQPASSL	198
AAV7/1-738	105	ADTDSFGGNGLRVFAQAKRRLLEPLGLVVEAAKTAAPKRRPVEQSPQEPDSSASISGKTSSTSSDAEAGPSGSLQLQIAQPASSL	198
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AAV36/1-737	105	ADTDSFGGNGLRVFAQAKRRLLEPLGLVVEAAKTAAPKRRPVEQSPQEPDSSASISGKTSSTSSDAEAGPSGSLQLQIAQPASSL	198
AAV/LK03/1-737	105	ADTDSFGGNGLRVFAQAKRRLLEPLGLVVEAAKTAAPKRRPVEQSPQEPDSSASISGKTSSTSSDAEAGPSGSLQLQIAQPASSL	198
AAV8/1-739	105	ADTDSFGGNGLRVFAQAKRRLLEPLGLVVEAAKTAAPKRRPVEQSPQEPDSSASISGKTSSTSSDAEAGPSGSLQLQIAQPASSL	199
AAV10/1-739	105	ADTDSFGGNGLRVFAQAKRRLLEPLGLVVEAAKTAAPKRRPVEQSPQEPDSSASISGKTSSTSSDAEAGPSGSLQLQIAQPASSL	199
AAV2/1-736	105	ADTDSFGGNGLRVFAQAKRRLLEPLGLVVEAAKTAAPKRRPVEQSPQEPDSSASISGKTSSTSSDAEAGPSGSLQLQIAQPASSL	198
AAV/DJ/1-738	105	ADTDSFGGNGLRVFAQAKRRLLEPLGLVVEAAKTAAPKRRPVEQSPQEPDSSASISGKTSSTSSDAEAGPSGSLQLQIAQPASSL	198
<b>VR1</b>			
AAV5/1-725	180	ADTMSAGGGRLDNNQAGADGVGNSAGDWHDCSTWMDRVTIKSTRVWVLSYNNHQRREIKGGSV-DGSSNANAYEGYSTPWGYDFNRFHCHFSRDPWRQL	291
AAV12/1-743	202	HDAENRAAPPGNAVEAGGGADGVGNSAGDWHDCSTWMDRVTIKSTRVWVLSYNNHQRREIKGGSV-DGSSNANAYEGYSTPWGYDFNRFHCHFSRDPWRQL	300
AAV9/1-737	190	SLTIMASGGGAFVADNNEGADGVGNSAGDWHDCSTWMDRVTIKSTRVWVLSYNNHQRREIKGGSV-DGSSNANAYEGYSTPWGYDFNRFHCHFSRDPWRQL	302
AAV1/1-737	190	SLTIMASGGGAFVADNNEGADGVGNSAGDWHDCSTWMDRVTIKSTRVWVLSYNNHQRREIKGGSV-DGSSNANAYEGYSTPWGYDFNRFHCHFSRDPWRQL	301
AAV6/1-737	190	SLTIMASGGGAFVADNNEGADGVGNSAGDWHDCSTWMDRVTIKSTRVWVLSYNNHQRREIKGGSV-DGSSNANAYEGYSTPWGYDFNRFHCHFSRDPWRQL	301
AAV7/1-738	200	SGSTIAAGGGAFVADNNEGADGVGNSAGDWHDCSTWMDRVTIKSTRVWVLSYNNHQRREIKGGSV-DGSSNANAYEGYSTPWGYDFNRFHCHFSRDPWRQL	302
AAV13/1-734	190	SLTIMASGGGAFVADNNEGADGVGNSAGDWHDCSTWMDRVTIKSTRVWVLSYNNHQRREIKGGSV-DGSSNANAYEGYSTPWGYDFNRFHCHFSRDPWRQL	299
AAV36/1-737	190	SLTIMASGGGAFVADNNEGADGVGNSAGDWHDCSTWMDRVTIKSTRVWVLSYNNHQRREIKGGSV-DGSSNANAYEGYSTPWGYDFNRFHCHFSRDPWRQL	300
AAV/LK03/1-737	190	SLTIMASGGGAFVADNNEGADGVGNSAGDWHDCSTWMDRVTIKSTRVWVLSYNNHQRREIKGGSV-DGSSNANAYEGYSTPWGYDFNRFHCHFSRDPWRQL	300
AAV8/1-739	200	PNTMAAGGGAFVADNNEGADGVGNSAGDWHDCSTWMDRVTIKSTRVWVLSYNNHQRREIKGGSV-DGSSNANAYEGYSTPWGYDFNRFHCHFSRDPWRQL	303
AAV10/1-739	200	PNTMAAGGGAFVADNNEGADGVGNSAGDWHDCSTWMDRVTIKSTRVWVLSYNNHQRREIKGGSV-DGSSNANAYEGYSTPWGYDFNRFHCHFSRDPWRQL	303
AAV2/1-736	190	PNTMAAGGGAFVADNNEGADGVGNSAGDWHDCSTWMDRVTIKSTRVWVLSYNNHQRREIKGGSV-DGSSNANAYEGYSTPWGYDFNRFHCHFSRDPWRQL	300
AAV/DJ/1-738	190	SLTIMASGGGAFVADNNEGADGVGNSAGDWHDCSTWMDRVTIKSTRVWVLSYNNHQRREIKGGSV-DGSSNANAYEGYSTPWGYDFNRFHCHFSRDPWRQL	302
AAV5/1-725	292	NNYWGFRPKRLSFKLFIQVKEVTNDGVTIANNLSTVQVFDSDSEYQLPYLGSAGHGCLPPFADVFMIPQGYLTLNNGS-DAGVRSFSYCLEYFP	395
AAV12/1-743	301	NNNWLFRPKRMLKFLFIQVKEVTNSRETTIANNLSTVQVFDSDSEYQLPYLGSAGHGCLPPFADVFMIPQGYLTLNNGS-DAGVRSFSYCLEYFP	404
AAV9/1-737	301	NNNWFPRKRLNFKLFIQVKEVTNDGVTIANNLSTVQVFDSDSEYQLPYLGSAGHGCLPPFADVFMIPQGYLTLNNGS-DAGVRSFSYCLEYFP	404
AAV1/1-737	302	NNNWFPRKRLNFKLFIQVKEVTNDGVTIANNLSTVQVFDSDSEYQLPYLGSAGHGCLPPFADVFMIPQGYLTLNNGS-DAGVRSFSYCLEYFP	403
AAV6/1-737	302	NNNWFPRKRLNFKLFIQVKEVTNDGVTIANNLSTVQVFDSDSEYQLPYLGSAGHGCLPPFADVFMIPQGYLTLNNGS-DAGVRSFSYCLEYFP	403
AAV7/1-738	303	NNNWFPRKRLNFKLFIQVKEVTNDGVTIANNLSTVQVFDSDSEYQLPYLGSAGHGCLPPFADVFMIPQGYLTLNNGS-DAGVRSFSYCLEYFP	404
AAV13/1-734	300	NNNWFPRKRLNFKLFIQVKEVTNDGVTIANNLSTVQVFDSDSEYQLPYLGSAGHGCLPPFADVFMIPQGYLTLNNGS-DAGVRSFSYCLEYFP	401
AAV36/1-737	301	NNNWFPRKRLNFKLFIQVKEVTNDGVTIANNLSTVQVFDSDSEYQLPYLGSAGHGCLPPFADVFMIPQGYLTLNNGS-DAGVRSFSYCLEYFP	402
AAV/LK03/1-737	301	NNNWFPRKRLNFKLFIQVKEVTNDGVTIANNLSTVQVFDSDSEYQLPYLGSAGHGCLPPFADVFMIPQGYLTLNNGS-DAGVRSFSYCLEYFP	402
AAV8/1-739	304	NNNWFPRKRLNFKLFIQVKEVTNEBTKIANNLSTVQVFDSDSEYQLPYLGSAGHGCLPPFADVFMIPQGYLTLNNGS-DAGVRSFSYCLEYFP	405
AAV10/1-739	304	NNNWFPRKRLNFKLFIQVKEVTNEBTKIANNLSTVQVFDSDSEYQLPYLGSAGHGCLPPFADVFMIPQGYLTLNNGS-DAGVRSFSYCLEYFP	405
AAV2/1-736	301	NNNWFPRKRLNFKLFIQVKEVTNDGVTIANNLSTVQVFDSDSEYQLPYLGSAGHGCLPPFADVFMIPQGYLTLNNGS-DAGVRSFSYCLEYFP	402
AAV/DJ/1-738	303	NNNWFPRKRLNFKLFIQVKEVTNEBTKIANNLSTVQVFDSDSEYQLPYLGSAGHGCLPPFADVFMIPQGYLTLNNGS-DAGVRSFSYCLEYFP	404
<b>VRIV</b>			
AAV5/1-725	396	LRTGNFESYTFEDEVPHSSYAHQSGLDRLMNPLIDQYLYVRSNTTGSNLTGQATTTTGYKITTDGFAYRRKNWLPAGIKDQKFSKNNANQYKIPASGGDA	487
AAV12/1-743	405	LRTGNFESYTFEDEVPHSSYAHQSGLDRLMNPLIDQYLYVRSNTTGSNLTGQATTTTGYKITTDGFAYRRKNWLPAGIKDQKFSKNNANQYKIPASGGDA	508
AAV9/1-737	404	LRTGNFESYTFEDEVPHSSYAHQSGLDRLMNPLIDQYLYVRSNTTGSNLTGQATTTTGYKITTDGFAYRRKNWLPAGIKDQKFSKNNANQYKIPASGGDA	501
AAV1/1-737	404	LRTGNFESYTFEDEVPHSSYAHQSGLDRLMNPLIDQYLYVRSNTTGSNLTGQATTTTGYKITTDGFAYRRKNWLPAGIKDQKFSKNNANQYKIPASGGDA	501
AAV6/1-737	404	LRTGNFESYTFEDEVPHSSYAHQSGLDRLMNPLIDQYLYVRSNTTGSNLTGQATTTTGYKITTDGFAYRRKNWLPAGIKDQKFSKNNANQYKIPASGGDA	501
AAV7/1-738	405	LRTGNFESYTFEDEVPHSSYAHQSGLDRLMNPLIDQYLYVRSNTTGSNLTGQATTTTGYKITTDGFAYRRKNWLPAGIKDQKFSKNNANQYKIPASGGDA	503
AAV13/1-734	403	LRTGNFESYTFEDEVPHSSYAHQSGLDRLMNPLIDQYLYVRSNTTGSNLTGQATTTTGYKITTDGFAYRRKNWLPAGIKDQKFSKNNANQYKIPASGGDA	498
AAV36/1-737	403	LRTGNFESYTFEDEVPHSSYAHQSGLDRLMNPLIDQYLYVRSNTTGSNLTGQATTTTGYKITTDGFAYRRKNWLPAGIKDQKFSKNNANQYKIPASGGDA	501
AAV/LK03/1-737	403	LRTGNFESYTFEDEVPHSSYAHQSGLDRLMNPLIDQYLYVRSNTTGSNLTGQATTTTGYKITTDGFAYRRKNWLPAGIKDQKFSKNNANQYKIPASGGDA	501
AAV8/1-739	406	LRTGNFESYTFEDEVPHSSYAHQSGLDRLMNPLIDQYLYVRSNTTGSNLTGQATTTTGYKITTDGFAYRRKNWLPAGIKDQKFSKNNANQYKIPASGGDA	503
AAV10/1-739	406	LRTGNFESYTFEDEVPHSSYAHQSGLDRLMNPLIDQYLYVRSNTTGSNLTGQATTTTGYKITTDGFAYRRKNWLPAGIKDQKFSKNNANQYKIPASGGDA	503
AAV2/1-736	403	LRTGNFESYTFEDEVPHSSYAHQSGLDRLMNPLIDQYLYVRSNTTGSNLTGQATTTTGYKITTDGFAYRRKNWLPAGIKDQKFSKNNANQYKIPASGGDA	500
AAV/DJ/1-738	405	LRTGNFESYTFEDEVPHSSYAHQSGLDRLMNPLIDQYLYVRSNTTGSNLTGQATTTTGYKITTDGFAYRRKNWLPAGIKDQKFSKNNANQYKIPASGGDA	502
<b>VRV</b>			
AAV5/1-725	489	AFATNRMREIASGYQPPFNGMNTNLQGSNTYALENTLNSQANPGETTATYLGEMLLIISEETOPVNRVYIVGQMTANSSITAPAGTYMLI	591
AAV12/1-743	508	LLKYDTHHTNWRGNNAAPGPIATAGAGDSDNSGLIFAGPNPNSNTTSSNLLFISEETOPVNRVYIVGQMTANSSITAPAGTYMLI	608
AAV9/1-737	502	AWPQASXWAGNRSLVNPQAMASHKDEEKFFPMHNLIFQHGHTAASNAEDLNMIIDEEIRITNPVATQYGVSNLQSSNAPTIRTNDGAL	602
AAV1/1-737	502	TWTQASKYHNGREIINPQTAMASHKDEEKFFPMHNLIFQHGHTAASNAEDLNMIIDEEIRITNPVATQYGVSNLQSSNAPTIRTNDGAL	602
AAV6/1-737	502	TWTQASKYHNGREIINPQTAMASHKDEEKFFPMHNLIFQHGHTAASNAEDLNMIIDEEIRITNPVATQYGVSNLQSSNAPTIRTNDGAL	602
AAV7/1-738	504	AWTGAATKYHNGRSLVNPQAMASHKDEEKFFPMHNLIFQHGHTAASNAEDLNMIIDEEIRITNPVATQYGVSNLQSSNAPTIRTNDGAL	603
AAV13/1-734	500	PWTQASKYHNGRSLVNPQAMASHKDEEKFFPMHNLIFQHGHTAASNAEDLNMIIDEEIRITNPVATQYGVSNLQSSNAPTIRTNDGAL	599
AAV36/1-737	502	PWTQASKYHNGRSLVNPQAMASHKDEEKFFPMHNLIFQHGHTAASNAEDLNMIIDEEIRITNPVATQYGVSNLQSSNAPTIRTNDGAL	602
AAV/LK03/1-737	502	PWTQASKYHNGRSLVNPQAMASHKDEEKFFPMHNLIFQHGHTAASNAEDLNMIIDEEIRITNPVATQYGVSNLQSSNAPTIRTNDGAL	602
AAV8/1-739	504	AWTGAATKYHNGRSLVNPQAMASHKDEEKFFPMHNLIFQHGHTAASNAEDLNMIIDEEIRITNPVATQYGVSNLQSSNAPTIRTNDGAL	604
AAV10/1-739	504	AWTGAATKYHNGRSLVNPQAMASHKDEEKFFPMHNLIFQHGHTAASNAEDLNMIIDEEIRITNPVATQYGVSNLQSSNAPTIRTNDGAL	604
AAV2/1-736	501	AWTGAATKYHNGRSLVNPQAMASHKDEEKFFPMHNLIFQHGHTAASNAEDLNMIIDEEIRITNPVATQYGVSNLQSSNAPTIRTNDGAL	601
AAV/DJ/1-738	503	AWTGAATKYHNGRSLVNPQAMASHKDEEKFFPMHNLIFQHGHTAASNAEDLNMIIDEEIRITNPVATQYGVSNLQSSNAPTIRTNDGAL	603
<b>VRVIII</b>			
AAV5/1-725	592	PGMVWQRDQVYVQIWAQIPIHTDGHFHPSPLMGGFGLKHPPPOILIKNTVPANPPTTSPAAKFAFITYOYSTGQVSEIWELOKNSRWNPEIQYTSNYY	694
AAV12/1-743	603	PGMVWQRDQVYVQIWAQIPIHTDGHFHPSPLMGGFGLKHPPPOILIKNTVPANPPTTSPAAKFAFITYOYSTGQVSEIWELOKNSRWNPEIQYTSNYY	712
AAV9/1-737	603	PGMVWQRDQVYVQIWAQIPIHTDGHFHPSPLMGGFGLKHPPPOILIKNTVPANPPTTSPAAKFAFITYOYSTGQVSEIWELOKNSRWNPEIQYTSNYY	706
AAV1/1-737	603	PGMVWQRDQVYVQIWAQIPIHTDGHFHPSPLMGGFGLKHPPPOILIKNTVPANPPTTSPAAKFAFITYOYSTGQVSEIWELOKNSRWNPEIQYTSNYY	706
AAV6/1-737	603	PGMVWQRDQVYVQIWAQIPIHTDGHFHPSPLMGGFGLKHPPPOILIKNTVPANPPTTSPAAKFAFITYOYSTGQVSEIWELOKNSRWNPEIQYTSNYY	706
AAV7/1-738	604	PGMVWQRDQVYVQIWAQIPIHTDGHFHPSPLMGGFGLKHPPPOILIKNTVPANPPTTSPAAKFAFITYOYSTGQVSEIWELOKNSRWNPEIQYTSNYY	707
AAV13/1-734	600	PGMVWQRDQVYVQIWAQIPIHTDGHFHPSPLMGGFGLKHPPPOILIKNTVPANPPTTSPAAKFAFITYOYSTGQVSEIWELOKNSRWNPEIQYTSNYY	703
AAV36/1-737	603	PGMVWQRDQVYVQIWAQIPIHTDGHFHPSPLMGGFGLKHPPPOILIKNTVPANPPTTSPAAKFAFITYOYSTGQVSEIWELOKNSRWNPEIQYTSNYY	706
AAV/LK03/1-737	603	PGMVWQRDQVYVQIWAQIPIHTDGHFHPSPLMGGFGLKHPPPOILIKNTVPANPPTTSPAAKFAFITYOYSTGQVSEIWELOKNSRWNPEIQYTSNYY	706
AAV8/1-739	605	PGMVWQRDQVYVQIWAQIPIHTDGHFHPSPLMGGFGLKHPPPOILIKNTVPANPPTTSPAAKFAFITYOYSTGQVSEIWELOKNSRWNPEIQYTSNYY	708
AAV10/1-739	605	PGMVWQRDQVYVQIWAQIPIHTDGHFHPSPLMGGFGLKHPPPOILIKNTVPANPPTTSPAAKFAFITYOYSTGQVSEIWELOKNSRWNPEIQYTSNYY	708
AAV2/1-736	602	PGMVWQRDQVYVQIWAQIPIHTDGHFHPSPLMGGFGLKHPPPOILIKNTVPANPPTTSPAAKFAFITYOYSTGQVSEIWELOKNSRWNPEIQYTSNYY	705
AAV/DJ/1-738	604	PGMVWQRDQVYVQIWAQIPIHTDGHFHPSPLMGGFGLKHPPPOILIKNTVPANPPTTSPAAKFAFITYOYSTGQVSEIWELOKNSRWNPEIQYTSNYY	707
AAV5/1-725	695	DPQFVDEAPDSTGEVTRTRRIGRYLTRLNPL	724
AAV12/1-743	713	TQNSMLWAFANWAGNHELRARIGSRLTHHL	742
AAV9/1-737	707	KSNVDFVDTNGVYSEPRRIGRYLTRLNPL	736
AAV1/1-737	707	KSNVDFVDTNGVYSEPRRIGRYLTRLNPL	736
AAV6/1-737	707	KSNVDFVDTNGVYSEPRRIGRYLTRLNPL	736
AAV7/1-738	708	KSNVDFVDTNGVYSEPRRIGRYLTRLNPL	737
AAV13/1-734	704	KSNVDFVDTNGVYSEPRRIGRYLTRLNPL	733
AAV36/1-737	707	KSNVDFVDTNGVYSEPRRIGRYLTRLNPL	736
AAV/LK03/1-737	707	KSNVDFVDTNGVYSEPRRIGRYLTRLNPL	736
AAV8/1-739	709	KSNVDFVDTNGVYSEPRRIGRYLTRLNPL	738
AAV10/1-739	709	KSNVDFVDTNGVYSEPRRIGRYLTRLNPL	738
AAV2/1-736	706	KSNVDFVDTNGVYSEPRRIGRYLTRLNPL	735
AAV/DJ/1-738	708	KSNVDFVDTNGVYSEPRRIGRYLTRLNPL	737

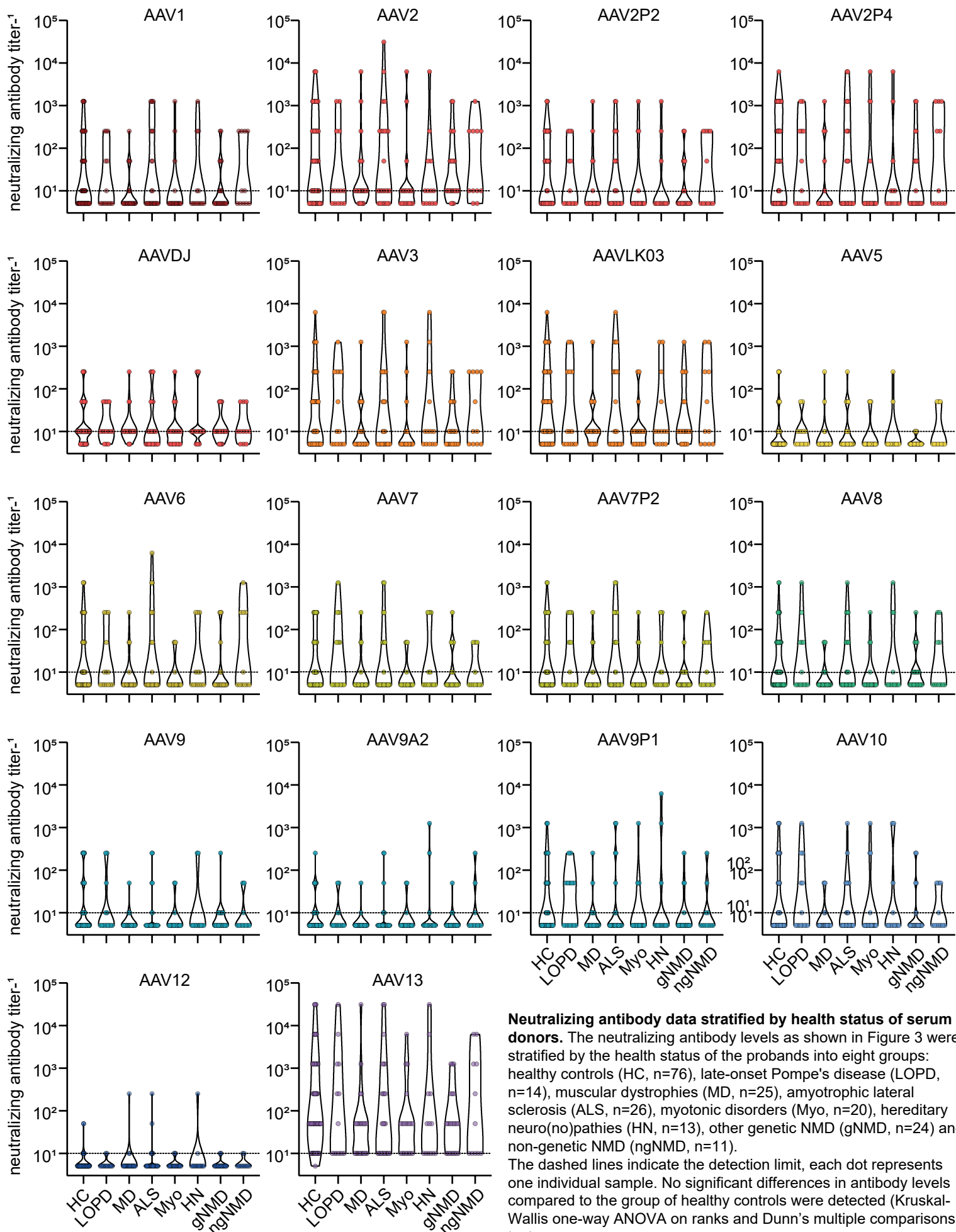
Multiple sequence alignment of wild-type and shuffled capsids used in this study. The alignments were generated using Clustal Omega 1.2.4, coloring is according to percentage identity and was generated in Jalview 2.11.3.3. Regions highlighted in light yellow, yellow, orange and red correspond to the variable regions I, IV, V and VIII, respectively. The green line represents the peptide insertion site in AAV2, 7 and 9.

Supplementary Figure S2

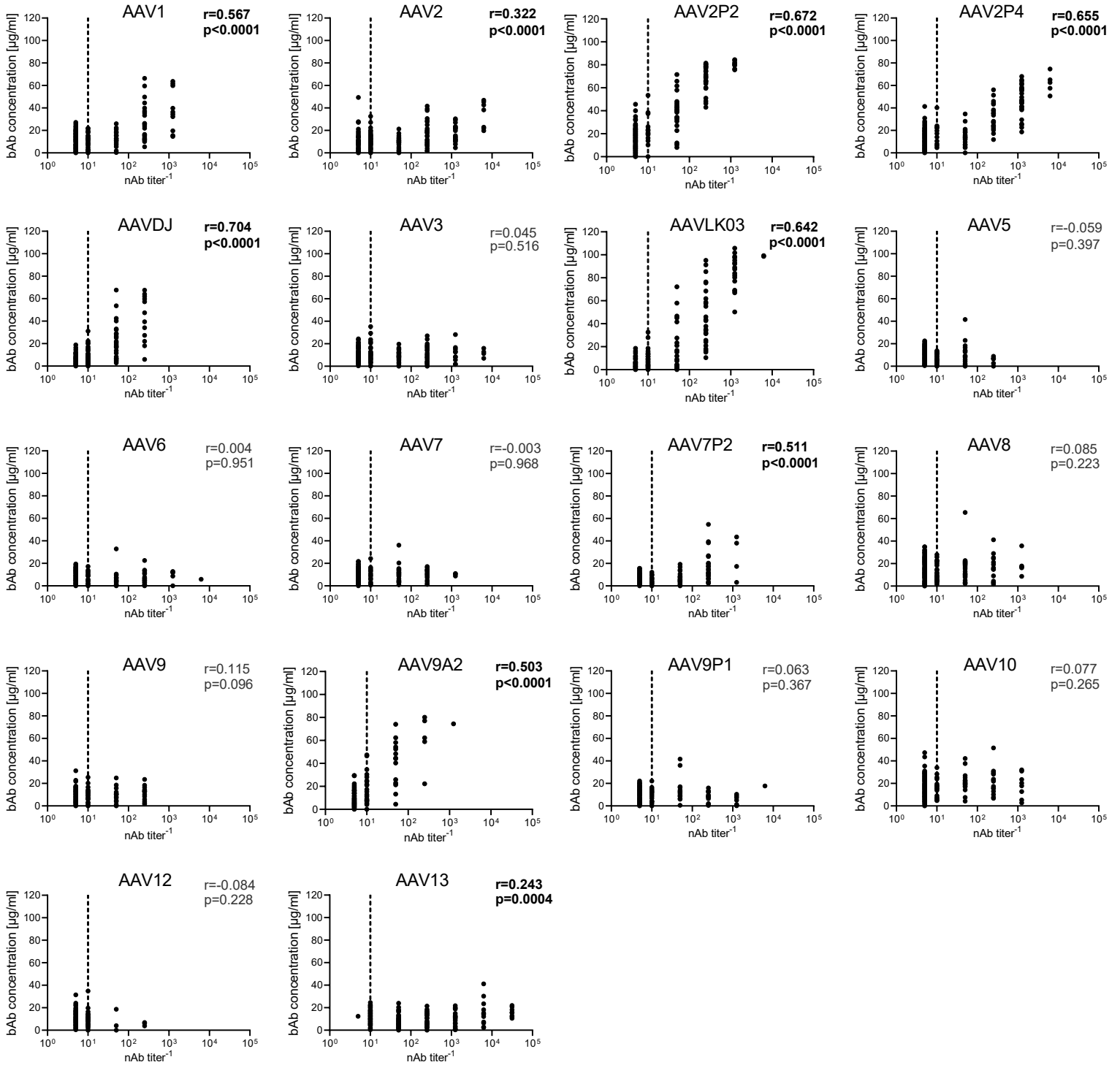


**Correlation of AAV-specific nAb levels and age of the serum donors.** The nAb titers against the indicated AAVs as shown in Figure 3 and the age of the serum donors were subjected to a Spearman ranked correlation analysis. Each dot indicates an individual donor, the dashed line indicates the nAb detection limit.

Supplementary Figure S3

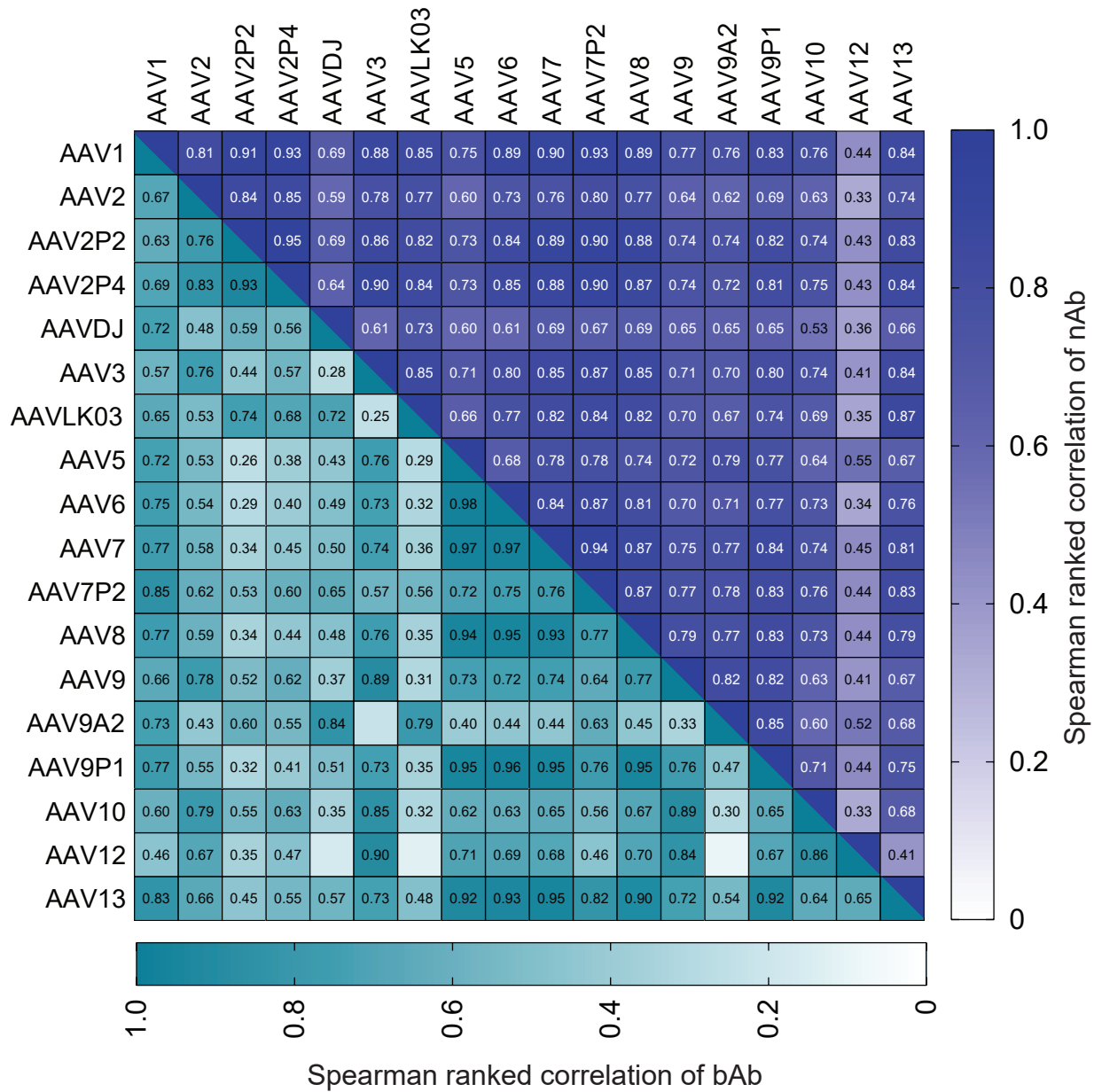


## Supplementary Figure S4



**Correlation of AAV-specific bAb and nAb levels.** The bAb titers as shown in Figure 2 and the nAb titers as shown in Figure 3 were subjected to a Spearman ranked correlation analysis.

Each dot indicates an individual donor, the dashed line indicates the nAb detection limit.



**Correlation between bAb and nAb levels specific for different AAV types.** A Spearman ranked correlation analysis was performed of the bAb levels as shown in Figure 2 (bottom left half-square) and of the nAb levels against the different AAV types as shown in Figure 3 (upper right half-square).

The colours indicate the correlation coefficient, values within the cells indicate the correlation coefficient if  $p < 0.001$ .