

Construct	MW (Da)	ϵ_{280} (M ⁻¹ cm ⁻¹)	Construct	MW (Da)	ϵ_{280} (M ⁻¹ cm ⁻¹)	Construct	MW (Da)	ϵ_{280} (M ⁻¹ cm ⁻¹)
<i>Pf</i> LDH	34930.6	17420	<i>Pf</i> MDH-R102K	35077.7	11920	AncMDH2-R102K-INS	36382.9	25900
<i>Pf</i> LDH_T101A	34900.5	17420	<i>Pf</i> MDH-INS	35751.3	17420	AncLDH	36969.6	27390
<i>Pf</i> LDH_K102A	34873.5	17420	<i>Cp</i> MDH-INS	35335.7	20400	AncMDH2-R102L	35725.3	20400
<i>Pf</i> LDH_A103S	34946.6	17420	<i>Pf</i> MDH-R102K-INS	35723.3	17420	AncMDH2-R102G	35669.2	20400
<i>Pf</i> LDH_P105A	34904.5	17420	<i>Pf</i> LDH-K102R-DEL	34312.9	11920	AncMDH2-R102Q	35740.3	20400
<i>Pf</i> LDH_G106A	34944.6	17420	<i>Pf</i> LDH-DEL	34284.9	11920	AncMDH2-R102W	35798.3	25900
<i>Pf</i> LDH_K107aA	34873.5	17420	<i>Tg</i> LDH2-DEL	35729.5	11920	AncMDH2*	35360.1	11920
<i>Pf</i> LDH_S107bA	34914.6	17420	<i>Pf</i> LDH-K102R	34958.6	17420	AncMDH2*-INS	36002.8	17420
<i>Pf</i> LDH_D107cA	34886.5	17420	<i>Tg</i> LDH2	36372.2	17420	AncMDH2*-58Mut-R102K	35964.8	11920
<i>Pf</i> LDH_K107dA	34873.5	17420	AncMDH2	35768.3	20400	AncLDH*	36607.4	17420
<i>Pf</i> LDH_E107eA	34872.5	17420	AncMDH2-59Mut	36355.0	21890	AncMDH3	35285.0	11920
<i>Pf</i> LDH_W107fA	34815.4	11920	AncMDH2-R102K	35740.3	20400	AncMDH1	35188.6	21890
<i>Pf</i> LDH_N108A	34887.5	17420	AncMDH2-R102K-59Mut	36327.0	21890	<i>Tg</i> LDH1	36613.5	23380
<i>Pf</i> MDH	35105.7	11920	AncMDH2-INS	36411.0	25900	<i>Rb</i> MDH	34716.3	14900
<i>Cp</i> MDH	34693.1	14900	AncMDH2-INS-59Mut	36997.6	27390			