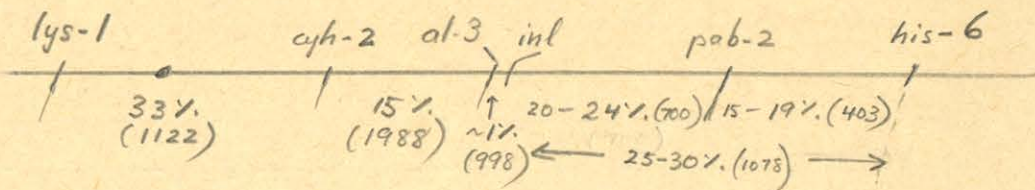


Mapping Strains (ctd)

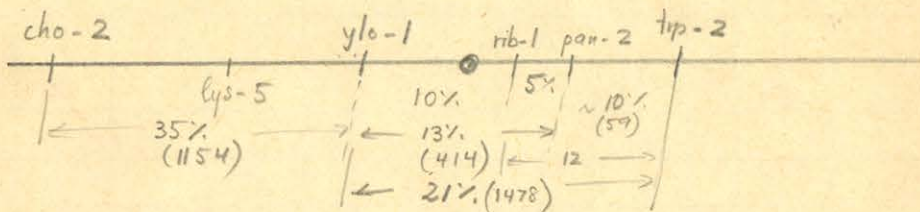
w M silica
 FGSC gel No |
 #

Linkage group V



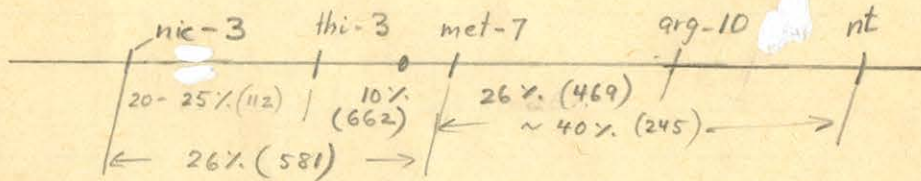
4129	1291 a	<i>lys-1</i>	<i>cyh-2</i>	<i>al-3</i>	<i>int</i> ^t	<i>pab-2</i>	<i>his-6</i>
4130	1287 A	<i>lys-1</i>	<i>cyh-2</i>	<i>al-3</i>		<i>pab-2</i>	<i>his-6</i>
4131	1009 A	<i>lys-1</i>	-	<i>al-3</i>	<i>int</i> ^t	<i>pab-2</i>	
4132	1008 a						
4133	1011 A	-	<i>cyh-2</i>	<i>al-3</i>	<i>int</i> ^t	<i>pab-2</i>	
4134	1010 a						
4135	to count 653 A	<i>lys-1</i>	-		<i>int</i>		<i>his-6</i>
4136	652 a						

VI



4137	854 A	<i>cho-2</i>	<i>ylo-1</i>			<i>tip-2</i> (15001)	} also doubles?
4138	1237 a						
4139	1806 A	-		<i>rib-1</i>	<i>pan-2</i> ^{B5}	<i>tip-2</i> (41)	} <i>rib-1 pan-2</i> (B2)
4140	1805 a						
							} <i>ylo-1 pan-2</i> (B3)

VII



4141	1208 A	<i>nic-3</i>	-	<i>met-7</i>	-	<i>arg-10</i>			
4142	1207 a								
4143	670 A	-		<i>thi-3</i>	<i>met-7</i>	-	<i>nt</i>		
4144	669 a								
4145	La. III	Translocation strain		11418	<i>acr-2</i>	<i>tip-1</i>	<i>ylo-1</i>	T(II,VI)	dow