	Core				Phenotyping			
	Generation/ Collection	Storage	Visualization	Analysis	Generation/ Collection	Storage	Visualization	Analysis
Field Book					<b>✓</b>	<b>√</b>		
GridScore					<b>✓</b>	<b>√</b>	<b>√</b>	
ClimMob	<b>✓</b>	<b>√</b>			<b>✓</b>	<b>√</b>		
ImageBreed						<b>√</b>	<b>√</b>	<b>✓</b>
PHIS		<b>√</b>			<b>✓</b>	<b>√</b>		<b>✓</b>
PIPPA		<b>√</b>			<b>✓</b>	<b>✓</b>		
Trait Selector BrAPP							<b>✓</b>	
Flapjack							<b>√</b>	
DArTView								
DivBrowse								
GIGWA								
PHG								
DArT Sample Submission								
MGIS								

AGENT Portal							
Helium					<b>√</b>	<b>✓</b>	
GLIS							
FLORILÈGE							
FAIDARE			✓				
DeltaBreed	<b>√</b>	<b>√</b>	<b>✓</b>		<b>√</b>	<b>✓</b>	<b>✓</b>
BMS	<b>√</b>	<b>√</b>	<b>✓</b>		<b>√</b>	<b>√</b>	<b>✓</b>
Breedbase	<b>√</b>	<b>√</b>	<b>✓</b>		<b>√</b>	<b>✓</b>	<b>✓</b>
BIMS	<b>√</b>	<b>√</b>	<b>✓</b>		<b>√</b>	<b>✓</b>	
Germinate	<b>√</b>	<b>√</b>	<b>✓</b>		<b>√</b>	<b>✓</b>	
QBMS				<b>√</b>			<b>✓</b>
Mr. Bean			<b>√</b>	<b>√</b>		<b>✓</b>	<b>✓</b>
G-Crunch				<b>√</b>			<b>✓</b>
SCT							

	Geno	typing		Germplasm					
Generation/ Collection	Storage	Visualization	Analysis	Generation/ Collection	Storage	Visualization	Analysis		
					<b>√</b>				
					<b>√</b>				
		<b>√</b>	<b>√</b>						
		<b>✓</b>	<b>√</b>						
	<b>√</b>	<b>√</b>	<b>√</b>						
	<b>√</b>		<b>√</b>						
<b>√</b>	<b>√</b>		<b>√</b>						
<b>√</b>									
				<b>✓</b>	<b>√</b>		<b>√</b>		

			<b>√</b>	<b>√</b>	<b>√</b>	
				<b>✓</b>	<b>√</b>	
			<b>✓</b>	<b>✓</b>		
				<b>✓</b>	<b>✓</b>	
			<b>✓</b>		<b>✓</b>	
<b>✓</b>				<b>✓</b>	<b>✓</b>	
<b>✓</b>				<b>✓</b>	<b>√</b>	
<b>√</b>				<b>√</b>	<b>√</b>	
				<b>√</b>	<b>√</b>	
<b>✓</b>				<b>✓</b>	<b>√</b>	
		<b>√</b>				<b>√</b>
	<b>✓</b>	<b>√</b>			<b>√</b>	<b>√</b>
		<b>✓</b>				<b>√</b>
					<b>√</b>	<b>√</b>