

# **Data Sheet**

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# pLSG-IBA123

Cat. No.: 5-4923-001 Lot No.: 4923-

Description

Affinity tag

0	)1	Version: Revision Date:	2.2 09.03.2020
	<ul> <li>StarGate® Acceptor Vector designed for gene transfer into the AcMNPV DNA by homologous recombination containing the for Polyhedrin promoter for high-level expression in inset.</li> <li>Co-transfection with BacPAK6 linearized AcMNPV DN circular <i>flash</i>BAC modified AcMNPV DNA (Oxford Exp allows the generation of recombinant baculovirus at through reconstitution of an essential gene (ORF 162 type virus to great extent.</li> <li>Ampicillin resistance and ColE1 origin of replication (<i>E. coli</i>.</li> <li>The expressed recombinant protein will be localized in the set of the</li></ul>	ollowing elen ct cells. A (Clontech) ression Tech very high effi 9) and elimin oUC) for prop	or with nologies) ciency ation of wild- pagation in
	The recombinant protein will contain two affinity tags:		

1.	GST-tag (Glutathione-S-Transferase) for the purification of recombinant protein.
	The affinity tag is fused to the N-terminus of the recombinant protein. After
	purification the GST may be removed by digesting with PreScission™ Protease.
2.	Strep-Tactin affinity tag (Twin-Strep-tag) for purification of recombinant protein
	via Strep-Tactin resin. The Twin-Strep-tag is fused to the C-terminus of the

	recombinant protein.
Resistance	Ampicillin
Form	5 μg, dissolved in 20 μl TE buffer, pH 8,0: 10 mM Tris-HCl, 1 mM EDTA
Concentration	250 ng/μl
Stability	12 months after shipping
Storage	recommended: 2-8 °C for frequent usage, -20 °C for long-term storage
Shipping	room temperature
Hazards	Product is not classified as hazardous according to (EC) No 1272/2008 [CLP]. A Material Safety Data Sheet is provided.

Note: The sequences have been compiled from information in the sequence database, published literature, and other sources, together with partial sequences obtained by IBA, however, the vectors have not been completely sequenced.

#### For research use only

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35 IG 1 55 GG	R 136 GAC D 166 AGC S 196 AGC AGC	L 137 CCCC2 P 167 167 CCAA( K 197 197	C 138 AATCO M 168 GGAA 198 GGAA E E 228	H 139 GTGC C 169 CAT/ 199 199 AGT: V 229	K 140 CCTC L 170 AGC <i>I</i> 200 FCTC CS L 230	T 141 GGA D 171 171 4TGC 201 GTTC	Y 142 GGCC A 172 GGCC 172 202 CCAC	L 143 GTTC F 173 CTTC 203 GGGC G G 1	$\begin{array}{c} N \\ I \\ $	$\frac{G}{145}$ $\frac{K}{175}$ $\frac{K}{205}$ $\frac{G}{55p}$ $\frac{G}{3}$ $\frac{M}{3}$	D 146 ATTP 176 CTGC G W 206 I JAGA	H 147 AGTJ V 177 GCA <i>A</i> ST 207	V 148 77TG C 178 AGCC A 208	G T 149 FTTTT F T 179 CACG CACG	ST H 150 CAAZ ST ST 180 GTT1 210 integ	P 151 AAAAA K 181 CGGT 211 grati	D 152 ACGI R 182 CGGI g 1 212 on s	F 153 CATT 1 183 CGGC G G 213 Site	M 154 FGAA E 184 CCGAC D 214	L 155 AGCI A 185 CCAI H 1215	Y 156 PATC 216	D 157 0CCC2 P P 187 187 217 217		L 159 MATT 189 MATC 219 Esp Aggg	D 160 FGA D 190 GTC S 220 31 S 2 2	V 161 TTAA( 191 CCGG, P G T 221 CCGC' Kwin-S A 3	V 162 GTA( Y 192 AGGC CS G T 222 FTG( trep-t W 4	L 163 CTTC L 193 IGG( G C 223 G G 223 G G 5 S 5	
35 IG 1 55 GG	R 136 GAC D 166 AGC S 196 AGC AGC	L 137 CCCC2 P 167 167 CCAA( K 197 197	C 138 AATCO M 168 GGAA 198 GGAA E E 228	H 139 GTGC C 169 CAT/ 199 199 AGT: V 229	K 140 CCTC L 170 AGC <i>I</i> 200 FCTC CS L 230	T 141 GGA D 171 171 4TGC 201 GTTC	Y 142 GGCC A 172 GGCC 172 202 CCAC	L 143 GTTC F 173 CTTC 203 GGGC G G 1	$\begin{array}{c} N \\ I \\ $	$\frac{G}{145}$ $\frac{K}{175}$ $\frac{K}{205}$ $\frac{G}{55p}$ $\frac{G}{3}$ $\frac{M}{3}$	D 146 ATT77 176 CTGC G W 206 I JAG7	H 147 AGTJ V 177 GCA <i>A</i> ST 207	V 148 TTGT C 178 AGCC A 208 CG	G T 149 FTTTT G F T 179 209 209	ST H 150 CAAZ ST ST 180 GTT1 210 integ	P 151 AAAAA K 181 CGGT 211 grati	D 152 ACGI R 182 CGGI g 1 212 on s	F 153 CATT 1 183 CGGC G G 213 Site	M 154 FGAA E 184 CCGAC D 214	L 155 AGCI A 185 CCAI H 1215	Y 156 PATC 216	D 157 200022 P 187 217 217		L 159 MATJ 189 MATC 219 Esp Aggg	D 160 FGA D 190 GTC S 220 31 S 2 2	V 161 TTAA( 191 CCGG, P G T 221 CCGC' Kwin-S A 3	V 162 GTA( Y 192 AGGC CS G T 222 FTG( trep-t W 4	L 163 CTTC L 193 IGG( G C 223 G G 223 G G 5 S 5	
GG GG	R 136 GAC D 166 AGC S 196 AGC AGC	L 137 CCCC2 P 167 167 CCAA( K 197 197	C 138 AATCO M 168 GGAA 198 GGAA E E 228	H 139 GTGC C 169 CAT/ 199 199 AGT: V 229	K 140 CCTC L 170 AGC <i>I</i> 200 FCTC CS L 230	T 141 GGA D 171 171 4TGC 201 GTTC	Y 142 GGCC A 172 GGCC 172 202 CCAC	L 143 GTTC F 173 CTTC 203 GGGC G G 1	$\begin{array}{c} N \\ I \\ $	$\frac{G}{145}$ $\frac{K}{175}$ $\frac{K}{205}$ $\frac{G}{55p}$ $\frac{G}{3}$ $\frac{M}{3}$	D 146 ATT77 176 CTGC G W 206 I JAG7	H 147 AGTJ V 177 SCAZ ST Q 207 AGAC	V 148 TTGT C 178 AGCC A 208 CG	G T 149 FTTTT G F T 179 209 209	ST H 150 CAAZ ST ST 180 GTT1 210 integ	P 151 AAAAA K 181 CGGT 211 grati	D 152 ACGI R 182 CGGI g 1 212 on s	F 153 CATT 1 183 CGGC G G 213 Site	M 154 FGAA E 184 CCGAC D 214	L 155 AGCI A 185 CCAI H 1215	Y 156 PATC 216	D 157 200022 P 187 217 217		L 159 MATJ 189 MATC 219 Esp Aggg	D 160 FGA D 190 GTC S 220 31 S 2 2	V 161 TTAA( 191 CCGG, P G T 221 CCGC' Kwin-S A 3	V 162 GTA( Y 192 AGGC CS G T 222 FTG( trep-t W 4	L 163 CTTC L 193 IGG( G G 223 G G 223 G 3 223 S 5	
GG GG	R 1136 GAC D 1166 AGC S 196 AGC 226 CAG	L 137 CCC22 P 167 CAAC K 197 CCTC CCTC L 227 STTC	C 138 AATC M 168 GTA1 198 GGAZ E 228 CGAZ	H 139 GTGC C 169 CATA I 199 AGT 229	K 140 CCTC L 170 AGC <i>I</i> 200 FCTC CS L 230 AGG5	T 141 GGA1 D 171 ATG0 W 201 GTT0 F F 231	Y 142 GGCC A 172 GGCC 172 202 CCAC Q 232	L 143 GTTC F 173 CTTC 203 GGGC G G 1 1	P + 144 $P + 174$	G 145 AAAAA 175 GGGC G GGGC CGG1 CGG1	D 146 ATT77 176 CTGC G W 206 I JAG7 JAG7	H 147 AGTJ V 177 SCAZ ST Q 207 AGAC	V 148 C 178 AGCC A 208 CG	G: T 149 FTTTT CG F T 179 CACCO T 209 ; ; ; ; ; ; ; ; ; ; ; ; ; ; ; ; ; ; ;	ST H 150 CAAJ ST K 180 STT1 210 integ	P 151 AAAAA K 181 CGGT 211 grati	D 152 ACGI R 182 CGGI 212 ON S	F 153 CATT 1 183 CGGC <u>G</u> 213 site tein	M 154 FGAZ E 184 CGAC D 214	AGCI A 155 AGCI A 185 CCAI H 215	Y 156 CATC 1 186 2CC1 216	D 157 20002 P 187 217 217 CGT0 CGT0		L 159 MATT 189 MATC 219 Esp Aggg	D 160 FGA D 190 GTC S 220 31 S 2 2	V 161 TTAA( 191 CCGG, P G T 221 CCGC' Kwin-S A 3	V 162 GTA( Y 192 AGGC CS G T 222 FTG( trep-t W 4	L 163 CTTC L 193 IGG( G G 223 G G 223 G 3 223 S 5	

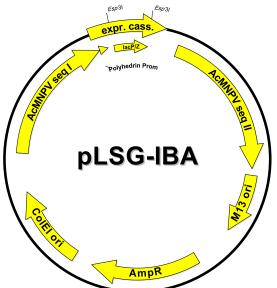
### Expression cassette of pLSG-IBA123, continued

LacP/Z cassette = contains LacZ alpha fragment under control of

of interest will be located here

a separate promoter, which allows alpha

complementation of *LacZ* mutations such as *LacZ*  $\Delta M15$  as in *E. coli* DH5 $\alpha$  or TOP10. after StarGate cloning using *Esp*3I your gene



Features Sequencing primer from bp to bp AcMNPVseq II 1 1395 LSG-Primer-for (Cat. No. 5-0000-161) M13 ori 1447 1920 5' - TAACCATCTCGCAAATAAATAAG -3' Ampicillin resistance gene 2251 3111 ColEI ori 3259 3902 AcMNPVseq I 4211 5357 LSG-Primer-rev (Cat. No. 5-0000-162) Polyhedrin promoter 5286 5355 forward primer binding site 5286 5308 5' - CAACGCACAGAATCTAGCGC -3' GST-tag 5369 6070 LacZ alpha fragment 6299 6700

6856

6941

6941

6764

6922



Twin-Strep-tag

total vector length

reverse primer binding site

your protein =

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