

Supplementary Information

Isolation, expression and biochemical characterization of recombinant hyoscyamine-6 β -hydroxylase from *Brugmansia sanguinea* – tuning the scopolamine production

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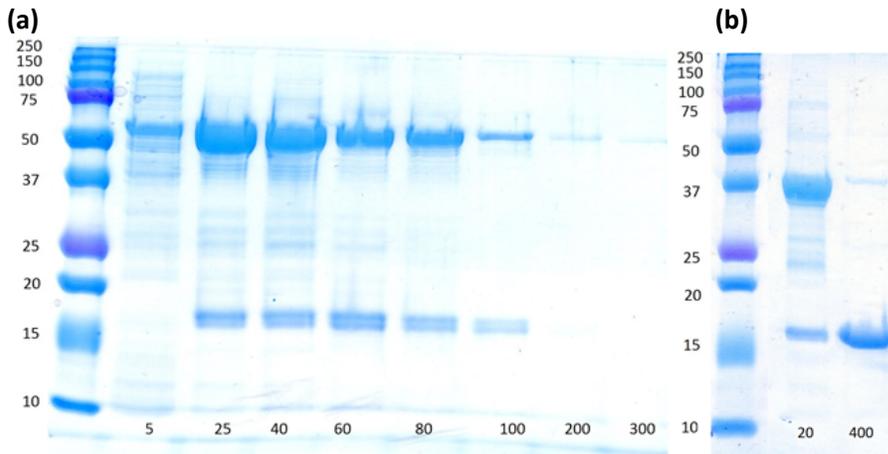


Figure S1. (a) Purification of SUMO-BsH6H analyzed by coomassie blue stained SDS-page. Amounts of imidazole used to elute the protein are indicated under each line (mM). (b) After cleavage of the SUMO-tag, the native BsH6H remains in the flowthrough (first lane).

		Fe-binding	
<i>Hyoscyamus niger</i>	MATFVSNWSTKSVSESFIAPIQKRAEKDVPIGNDVPIIDLQHHHLLVQQITKACQDFGL	60	
<i>Hyoscyamus senecionis</i>	MATFVSNWSTKSVTESFIAPIQKRAEKDVPIGNVPIIDLQHHHLLVQQITKACQDFGL	60	
<i>Datura stramonium</i>	MATFVSNWSTNNVSESFIAPLEKRAEKDVPLGNDVPIIDLQDHLIVQQITKACQDFGL	60	
<i>Datura metel</i>	MATLVSNWSTNNVSESFIAPLEKRAEKDVPLGNDVPIIDLQDHLIVQQITKACQDFGL	60	
<i>Duboisia myoporoides</i>	-----DVPIIDLKQDHIIVVQQINKACQDFGL	27	
<i>Brugmansia sanguinea</i>	MATFVSNWSTNNVSESFIAPLEKRAEKYVPLGNDVPIIDLQDHLIVQQITEACQDFGL	60	
<i>Brugmansia arborea</i>	MATFVSNWSTNNVSESFIAPLEKRAEKDVPLGNDVPIIDLQDHLIVQQITKACQDFGL	60	
<i>Atropa baetica</i>	MATFVSNWSTNNVSESFVAPLEKRAEKDVPLGNDVPTIDLQDHLVVVQQITKACQDFGL	60	
<i>Atropa belladonna</i>	MATLVSNWSTNNVSESFVAPLEKRAEKDVPLGNDVPIIDLQDHLVVVQQITKACQDFGL	60	
<i>Scopolia parviflora</i>	MATLVSNWSSNNVSESFIAPLEKRAEKDVPLGNDVPMIDLQDHHLLVQQITKACQDFGL	60	
<i>Anisodus acutangulus</i>	MATLVSNWSTNNVSESFKAPLEKRAEKDVPLGNDVPIIDLQDHLVVVQQITKACQDFGL	60	
<i>Anisodus tanguticus</i>	MATLVSNWSSNNVSESFIAPLDKRAEKVPLGNDVPIIDLQDHHFVVQQITKACKDFGL	60	
<i>Anisodus luridus</i>	MATFVSNWSSNNVSESFIAPLDKRAEKVPLGNDVPIIDLQDHHFVVQQITKACKDFGL	60	
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	Fe-binding		
<i>Hyoscyamus niger</i>	FQVINHGFPPEELMLETMEVCKEFFALPAEEKEKFKPKGEAAKFELPLEQKAKLYVEGEQL	120	
<i>Hyoscyamus senecioni</i>	FQVINHGFPPEELMLETMEVCKEFFALPAEEKEKFKPKGEAAKFELPLEQKAKLYIEGEQL	120	
<i>Datura stramonium</i>	FQVINHGVPPEKLMVEAMEVYKEFFALPAEEKEKFKPKGEPAKFELPLEQKAKLYVEGERR	120	
<i>Datura metel</i>	FQVINHGVPPEKLMVEAMEVYKEFFALPAEEKEKFKPKGEPAKFELPLEQKAKLYVEGERR	120	
<i>Duboisia myoporoides</i>	FQVINHGFPPEKLMVEAMEVYKEFFALPAEEKEKLPKGEPAKFELPLEQKAKLYVEGEQL	87	
<i>Brugmansia sanguinea</i>	FQVINHGFPSEKLMETMEVCKEFFALPAEEKEKLPKGEPAKFELPLEQKAKLYVEGEQL	120	
<i>Brugmansia arborea</i>	FQVINHGFPSEKLMETMEACKVEFFALPAEEKEKLPKGEPAKFELPLEQKAKLYVEGEQL	120	
<i>Atropa baetica</i>	FQVINHGLPEKLMETMDVCKEFFALPAEEKEKLPKGEPAKFELPLEQKAKLYVEGEQL	120	
<i>Atropa belladonna</i>	FQVINHGLPEKLMETMDVCKEFFALPAEEKEKLPKGEPAKFELPLEQKAKLYVEGEQL	120	
<i>Scopolia parviflora</i>	FQVINHGFPPEKLMETMKVCKEFFALPAEEKEKLPKGEPAKFELPLEQKAKLYIEGEQL	120	
<i>Anisodus acutangulus</i>	FQVINHGFPPEKLMETMKVCKEFFALPAEEKEKLPKGGKPAKFELPLEQKAKLYIEGEQL	120	
<i>Anisodus tanguticus</i>	FQVINHGFPPEKLMVETMEVCKEFFALPAEENEKLPKGGKPAKFELPLEQKAKLYIEGEQL	120	
<i>Anisodus luridus</i>	FQVINHGFPENLMAETMKLCKEFFALPAEEKEKLPKGGKPAKFELPLEQKAKLYIEGEQL	120	
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<i>Hyoscyamus niger</i>	SNEEFlyWkDtlAhGChPlDQdLVnSwPEkPAkYREvVAKYsVEVRkLTMRMLDYICEGL	180
<i>Hyoscyamus senecionis</i>	SNEEFlyWkDtlAhGChPlDQdLVnSwPEkPAkYREvVAKYsVEVRkLTMRMLDYISEGL	180
<i>Datura stramonium</i>	CNEEFlyWkDtlAhGcYpLHEELlNSWPEkPpTYRdVIAkYsVEVRkLTMRILDYICEGL	180
<i>Datura metel</i>	CNEEFlyWkDtlAhGcYpLHEELlNSWPEkPpTYRdVIAkYsVEVRkLTMRILDYICEGL	180
<i>Duboisia myoporoides</i>	SNDEFyWkDtlAhGChPlHEdLVnSwPDKpStYRElMSkYsVEVRkLTMRILDYICEGL	147
<i>Brugmansia sanguinea</i>	SNEEFlyWkDtlAhGChPlDEELlNSWPEkPpTYRdVMakYsVEVRkLTMRILDYICEGL	180
<i>Brugmansia arborea</i>	SNDEFlyWkDtlAhGChPlDEELlNSWPEkPpTYRdVMakYsVEVRkLTMRILDYICEGL	180
<i>Atropa baetica</i>	SDEAFlyWkDtlAhGChPlDEELlNSWPEkPATYREvVAKYsVEVRkLTMRILDYICEGL	180
<i>Atropa belladonna</i>	SDEAFlyWkNtlAhGChPlDEELlNSWPEkPATYREvVAKYsVEVRkLTMRILDYICEGL	180
<i>Scopolia parviflora</i>	SNEEFyWkDtlAhGChPlDEElKlVnSwPEkPATYREvVAKYsVEVRkLTMRMLDYICEGL	180
<i>Anisodus acutangulus</i>	SNGELfYwKdTLAhGChPlDEELVnSwPEkPATYREvVskYsVEVRkLTMRMLDYICEGL	180
<i>Anisodus tanguticus</i>	SNEEFyWkDtlAhGChPlDEELlNSWPEkPATYREvAAkYsVEARkLTMRMLDYICEGL	180
<i>Anisodus luridus</i>	SNDEFyWkDtlAhGChPlDEELVnSwPEkPttYREvAAkYsVEARkLTMRMLDYICEGL	180
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	2-ketoglutarate binding	
<i>Hyoscyamus niger</i>	GLKlGYFDnELsQIQmMLtNyYpPCDPsStlGSGGHyDGNlITllQqDlPGLQQLlVkd	240
<i>Hyoscyamus senecionis</i>	GLKlGYFDnELsQIQmMLtNyYpPCDPtStlGSGGHyDGNlITllQqDlPGLQqFIVkd	240
<i>Datura stramonium</i>	GLKlGYFDnELtQIQmLLAnYpScDPsStlGSGGHyDGNlITllQqDlVGLQQLlVkd	240
<i>Datura metel</i>	GLKlGYFDnELtQIQmLLAnYpScDPsStlGSGGHyDGNlITllQqDlVGLQQLlVkd	240
<i>Duboisia myoporoides</i>	GLKlGYFDnELcQIQmMLtNyYpPCDPsStlGSGGHyDGNlITllQqDlPGLQQLlVkd	207
<i>Brugmansia sanguinea</i>	GLKlGYFDnELsQIQmLLtNyYpPCDPsStlGSGGHyDGNlITllQqDlPGLQQLlVkd	240
<i>Brugmansia arborea</i>	GLKlGYFDnELsQIQmLLtNyYpPCDPsStlGSGGHyDGNlITllQqDlPGLQQLlVkd	240
<i>Atropa baetica</i>	GLKlGYFDnELsQIQmMLtNyYpPCDPsStlGSGGHyDGNlITllQqNlPGLQqFIVeD	240
<i>Atropa belladonna</i>	GLKlGYFDnELsQIQmMLtNyYpPCDPsStlGSGGHyDGNlITllQqNlPGLQqL-IEE	239
<i>Scopolia parviflora</i>	ELKlGYFDnELsQIQmMLtNyYpPCDPsStlGSGGHyDGNlITllQqDlPGLQQLlVkd	240
<i>Anisodus acutangulus</i>	GLKlGYFDnELsQIQmMLAnYpPCDPsStlGSGAHyDGNvITllQqDlPGLQQLlVkd	240
<i>Anisodus tanguticus</i>	GLKlGYFDnELsQIQmMLtNyYpPCDPsStlGSGGHyDGNlITllQqDlPGLQQLlVkd	240
<i>Anisodus luridus</i>	GLKlGHFDnELsQIQmMLtNyYpPCDPsStlGSGGHyDGNlITllQqDlPGLQQLlVkd	240
	****:***** ***:*.***** ***:*.***** ***:*.*****:*. *****:***	
	Fe-binding	
<i>Hyoscyamus niger</i>	ATWIAVQPIPTAFvVnLGLtLkVITNEkFEGSIHRvVtDpTRDRVSIATlIGPDYSCTIE	300
<i>Hyoscyamus senecionis</i>	STWIAVQPIPTAFvVnLGLtLkVIsNEkFEGSIHRvVtDpTRDRVSIATlIGPDYSCTIE	300
<i>Datura stramonium</i>	DKWIAVEPIPTAFvVnLGLtLkVMSNEkFEGSIHRvVtHPTRNRISIGtLIGPDYSCTIE	300
<i>Datura metel</i>	DKWIAVEPIPTAFvVnLGLtLkVMSNEkFEGSIHRvVtHPIRNRISIGtLIGPDYSCTIE	300
<i>Duboisia myoporoides</i>	GNWIAVEPIPTAFvVnLGLtLkVITNEkLEGSiHRvVtNPTRDRVSIATlIGPDYSCTIE	267
<i>Brugmansia sanguinea</i>	ANWIAVEPIPTAFvVnLGLtLkVIsNEkFEGSIHRvVtNPtSDRVSIATlIGPDYSCTIE	300
<i>Brugmansia arborea</i>	ANWIAVEPIPTAFvVnLGLtLkVIsNEkFEGSIHRvVtNPTRDRVSIATlIGPDYSCTIE	300
<i>Atropa baetica</i>	AKWIAVEPIPTAFvVnLGLtLkVITNEkFEGSIHRvVtNPTRDRVSIATlIGPDYSCTIE	300
<i>Atropa belladonna</i>	AKWIAVEPIPTAFvVnLGLtLkVITNEkFEGSIHRvVtNPTRDRVSIATlIGPDYSCTIE	299
<i>Scopolia parviflora</i>	GNWIAVEPNPTAFvInLGLtLkVITNEkFEGSIHRvVADpTRDRVSIATlIGPDYSCTIE	300
<i>Anisodus acutangulus</i>	DNWIAVEPIPTAFvInLGLtLkVITNEkFEGSIHRvVtNPTRDRVSIATlIGPDYSCTIE	300
<i>Anisodus tanguticus</i>	DNWIAVEPIPTAFvInLGLtLkVITNEkFEGSIHRvVtDpTRDRVSIATlIGPDYSCTIE	300
<i>Anisodus luridus</i>	DNWIAVEPIPTAFvInLGLtLkVITNEkFEGSIHRvVtNPTRDRVtIATlIGPDYSCTIE	300
	.***** * *****:*****:*.*****:*****:*. *****:*****	
<i>Hyoscyamus niger</i>	PAKELlNQDNpPlykPYsYSEfADIYlSDkSDYDsgVkpYkINv	344
<i>Hyoscyamus senecionis</i>	PAKELlNQDNpPlykPYsYAEfADIYlSDkSDYDagVrpYkINv	344
<i>Datura stramonium</i>	PIKELlSQENpPlykPYpYAKfAEIYlSDkSDYDagVkpYkIN-	343
<i>Datura metel</i>	PIKELlSQENpPlykPYpYAEfAEIYlSDkSDYDagVkpYkIN-	343
<i>Duboisia myoporoides</i>	PAKELlSQDNpPlykPYsYAEfAEIYlSDkSGYDagVkpYkLN-	310
<i>Brugmansia sanguinea</i>	PAKELvNQENpPlykPYpYAEfAEIYlSDkSDfDagVkpYkINv	344
<i>Brugmansia arborea</i>	PAKELvSQENpPlykPYpYvEAEfAEIYlSDkSDYDAvKpYkIN-	343
<i>Atropa baetica</i>	PAKELlSQDNpPlykPYsYAEfAEIYlSDkSDYDagVkpYkINv	344
<i>Atropa belladonna</i>	PAKELlSQDNpPlykPYsYAEfAEIYlSDkSDYDagVkpYkINa	343
<i>Scopolia parviflora</i>	PGKELlSQDNpPlykPYpYtEAEfAEIYlSDkSGY-----	333
<i>Anisodus acutangulus</i>	PAKELlSQDNpPlykPYpYAEfAEIYlSDkSGYDagVkpYkINa	344
<i>Anisodus tanguticus</i>	PAKELlSQDNpPlykPYpYAEfAEIYlSDkSGYDagVkpYkINa	344
<i>Anisodus luridus</i>	PAKELlSQDNpPlykPYpYAEfAEIYlSDkSGYDagVkpYkINa	344
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Figure S2. Multiple sequence alignments for comparison of BsH6H with other reported H6H homologues. Two iron binding and one co-factor region are boxed. Red letters indicate different residues with regard to the consensus. Blue highlighted letters depict significantly different residues in *Brugmansia sanguinea*-H6H compared to other H6H homologues. Accession numbers: ABG89397.1 (*H. niger*); AFP99876.1 (*H. senecionis*); ALD59774.1 (*D. stramonium*); AAQ04302.1 (*D. metel*);

AQU12714.1 (*D. myoporoides*); ALD59773.1 (*B. arborea*); ABR15749.1 (*A. baetica*); AEM91979.1 (*A. belladonna*); AAY53932.1 (*S. parviflora*); ABM74185.1 (*A. acutangulus*); AAQ75700.1 (*A. tanguticus*); AGL76991.1 (*A. luridus*).

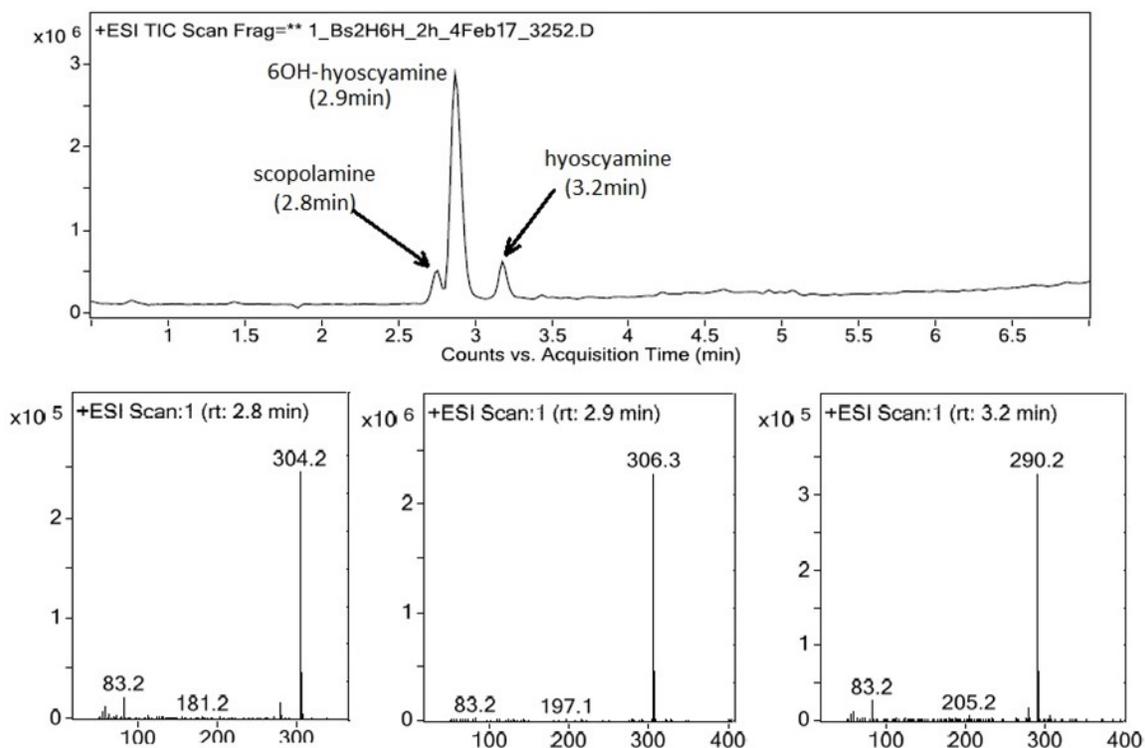


Figure S3. LCMS analysis of the transformation of hyoscyamine **1** (3.2 min) to intermediate **2** (2.9 min) and scopolamine **3** (2.8 min) by BsH6H and MS identification of products.