

Input	
Sequence ID	<input type="text" value="HBA_HORSE_P01958"/> ?
Sequence name	<input type="text" value="HBA_HORSE"/> ?
Sequence description	<input type="text"/> ?
Import sequence of	<input checked="" type="radio"/> aminoacids <input type="radio"/> nucleotides ?
From	<input type="radio"/> plain text <input type="radio"/> atomic structure <input type="radio"/> file <input checked="" type="radio"/> UniProt ID ?
UniProt name/ID	<input type="text" value="P01958"/> ?

Input	
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Sequence description	<input type="text"/> ?
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From	<input type="radio"/> plain text <input type="radio"/> atomic structure <input type="radio"/> file <input checked="" type="radio"/> UniProt ID ?
UniProt name/ID	<input type="text" value="P01948"/> ?

Input	
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Sequence description	<input type="text"/> ?
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From	<input type="radio"/> plain text <input type="radio"/> atomic structure <input type="radio"/> file <input checked="" type="radio"/> UniProt ID ?
UniProt name/ID	<input type="text" value="P81023"/> ?

Input	
Sequence ID	<input type="text" value="HBA_TURTLE_P83134"/> ?
Sequence name	<input type="text" value="HBA_TURTLE"/> ?
Sequence description	<input type="text"/> ?
Import sequence of	<input checked="" type="radio"/> aminoacids <input type="radio"/> nucleotides ?
From	<input type="radio"/> plain text <input type="radio"/> atomic structure <input type="radio"/> file <input checked="" type="radio"/> UniProt ID ?
UniProt name/ID	<input type="text" value="P83134"/> ?