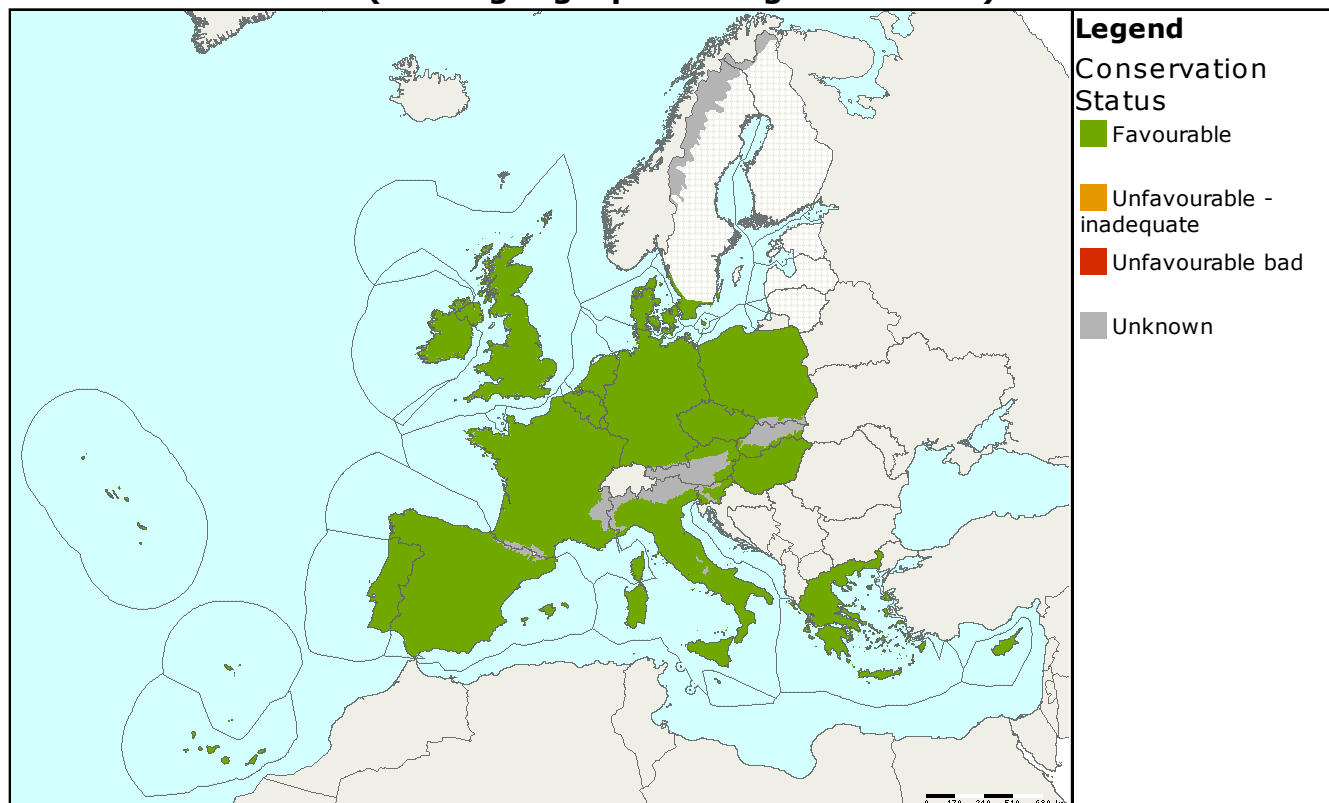


Species name: **Pipistrellus kuhlii**
Annex: **IV**

Species group: **Mammals**
Regions: **ALP ATL CON MAC MED PAN**

Assessments of conservation status at the European level (all biogeographical regions - EU25)



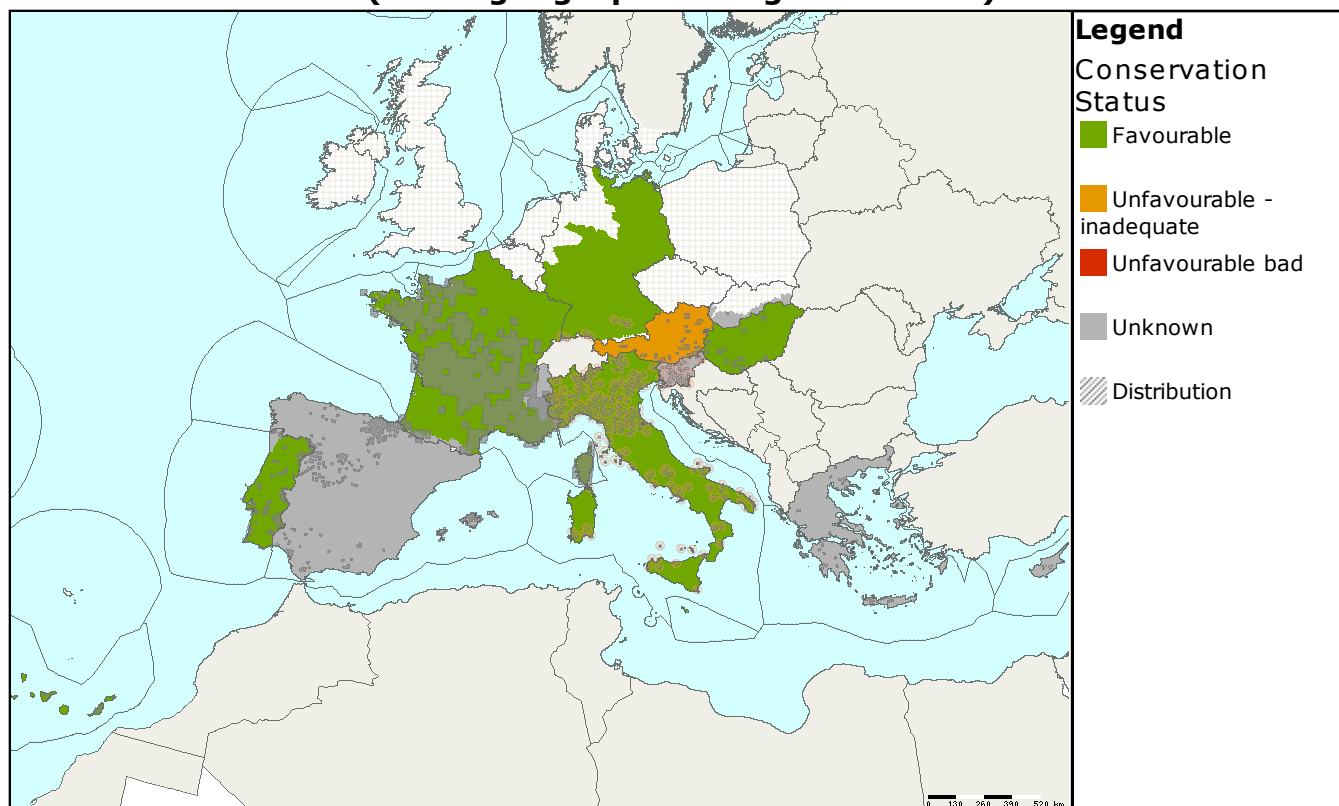
MS	Region	Conservation status assessment					Population size & unit	Population Trend
		Range	Population	Habitat	Future prospects	Overall		
EU25	ALP						440 grids	
EU25	ATL						1262 grids	
EU25	CON						1244 grids	
EU25	MAC							=
EU25	PAN						5000 - 30000 indiv.	+
EU25	MED						> 2004 grids	

Kuhl's pipistrelle is a bat with a large range extending from Iberia through southern Europe through the Near East and the Caucasus to Kazakhstan, Pakistan, and India. It forages over variety of habitats, including agricultural and urban areas.

Many countries reported individual parameters as 'unknown' showing that better information is required for this species. Overall assessment for five of the six regions where it occurs is 'favourable'. Its status is reported as 'unknown' in the Alpine region.

The species is widespread and abundant, and populations are increasing in parts of the range. Consequently it is assessed as 'Least Concern' by IUCN.

Assessments of conservation status as reported by Member states (all biogeographical regions - EU25)



MS	Region	Conservation status assessment					Size & unit	Population trend	Data quality
		Range	Population	Habitat	Future prospects	Overall			
AT	ALP	Unfavourable - inadequate	Unfavourable - inadequate	Favourable	Favourable	Unfavourable - inadequate	N/A x	+	3
ES	ALP	Unknown	Unknown	Unknown	Unknown	Unknown	0.5 - 0.5 loc.	X	3
FR	ALP	Favourable	Favourable	Favourable	Favourable	Favourable	N/A x	N/A	
IT	ALP	Favourable	Favourable	Favourable	Favourable	Favourable	74 - 74 loc.	=	2
SI	ALP	Favourable	Unknown	Unknown	Unknown	Unknown	N/A x	X	3
ES	ATL	Unknown	Unknown	Unknown	Unknown	Unknown	68 - 73 loc.	X	
FR	ATL	Favourable	Favourable	Favourable	Favourable	Favourable	N/A x	N/A	
PT	ATL	Favourable	Unknown	Unknown	Unknown	Unknown	N/A x	X	
AT	CON	Unfavourable - inadequate	Unfavourable - inadequate	Favourable	Favourable	Unfavourable - inadequate	N/A x	+	3
DE	CON	Favourable	Favourable	Favourable	Favourable	Favourable	10 - (10) x	+	2
FR	CON	Favourable	Unknown	Unknown	Unknown	Unknown	N/A x	N/A	
IT	CON	Favourable	Favourable	Favourable	Favourable	Favourable	242 - 242 loc.	=	2
SI	CON	Favourable	Unknown	Unknown	Unknown	Unknown	2 - (2) colony	X	3
ES	MAC	Unfavourable - inadequate	Favourable	Favourable	Favourable	Favourable	N/A x	=	
CY	MED	Unknown	Unknown	Unknown	Unknown	Unknown	4 - 4 grids	X	3
EL	MED	Unknown	Unknown	Unknown	Unknown	Unknown	64 - 64 x	X	3
ES	MED	Unknown	Unknown	Unknown	Unknown	Unknown	187 - 192 loc.	X	
FR	MED	Favourable	Favourable	Favourable	Favourable	Favourable	N/A x	N/A	
IT	MED	Favourable	Favourable	Favourable	Favourable	Favourable	381 - 381 loc.	=	2
MT	MED	Unknown	Unknown	Unknown	Unknown	Unknown	(500) - 500 indiv.	=	2
PT	MED	Favourable	Unknown	Unknown	Unknown	Unknown	10000 - (10000) indiv.	X	2
HU	PAN	Favourable	Favourable	Favourable	Favourable	Favourable	5000 - 30000 indiv.	+	3
SK	PAN	Unknown	Unknown	Unknown	Unknown	Unknown	1 - 1 indiv.	+	1

Data quality is based on an assessment by each Member State, 1 = good, 2 = medium,

3 = poor

This information is derived from the Member State national reports submitted to the European Commission under Article 17 of the Habitats Directive in 2007 and covering the period 2001-2006. More detailed information is available at <http://biodiversity.eionet.europa.eu/article17>