# **Spiders and venom**



M A S A R Y K U N I V E R S I T Y

### **Ondřej Michálek**



# OUTLINE

- 1) Introduction to venom & venom in spiders
- 2) Venom apparatus morphology
- 3) Venom chemistry
- 4) Venom evolution
- 5) Venom ecology
- 6) Methods to study venoms & venom applications
- 7) Sum-up

# Part 1: Introduction to venom

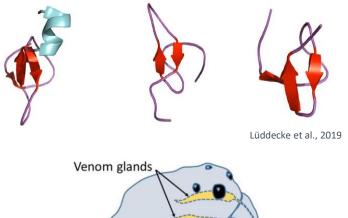


# What is venom?

- physical warfare -> chemical warfare
- "A biological substance produced by an organism that contains molecules ("toxins") which interfere with physiological or biochemical processes in another organism, which has evolved in the venomous organism to provide a benefit to itself once introduced to the other organism. The venom is produced and/or stored in a specialised structure and actively transferred to another organism through an injury by means of a specialised delivery system." (Arbuckle, 2017)

# What is venom?

- Key aspects of venom:
  - Toxins
  - Venom gland/tissue
  - Specialised venom apparatus
  - Transfer to target animal through injury
  - Alter physiological or biochemical processes in target animal (paralysis)
  - A benefit to venomous animal

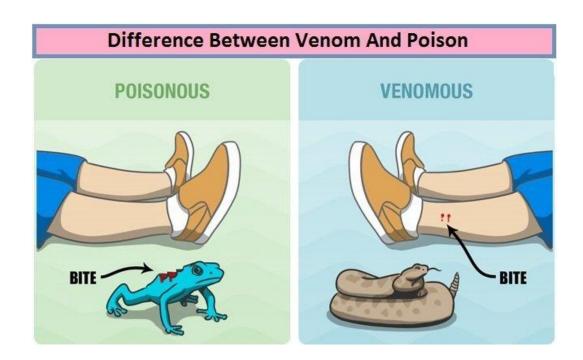








## Venom vs poison



#### toxic



birdandmoon.com

### Venom vs poison vs toxungen

Table 2. Critical components and features that distinguish the three major categories of biological toxins

Biological toxin	Delivery mechanism	Penetration wound	Mechanism of transfer or deployment
Poison	No	No	Ingestion, inhalation, or absorption across body surface
Toxungen	Yes	No	Delivered to body surface without accompanying wound
Venom	Yes	Yes	Delivered to internal tissues via wound



Photo: Wolfgang Wuster

Nelsen et al., 2014

## **Venom function**

- Predation
- Defence





## Venom function – not just predation and/or defence...

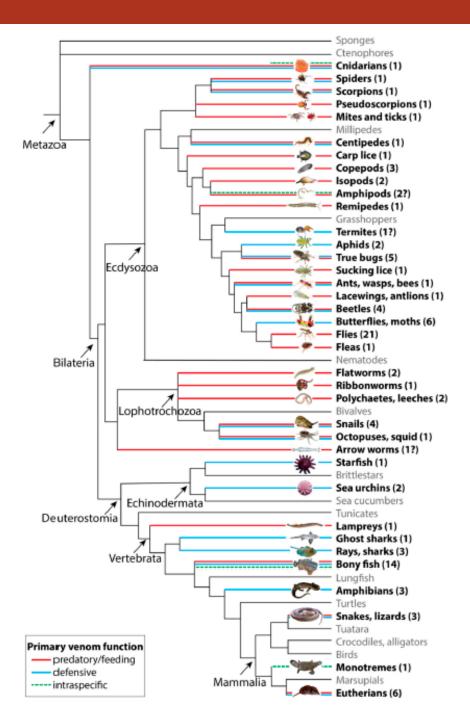
**Table 1.** Functional diversity of venom. Examples of uses of venom beyond predation, defense, and blood-feeding.

Function	<b>Example of Venomous Animal</b>	References
Intraspecific competition	Platypus, sea anemones, slow loris	[51-53]
Food storage	Moles, shrews, parasitoid wasps	[54,55]
(Pre-)Digestion	Sea anemones, assassin bugs, centipedes, remipedes, vipers	[56–60]
Offspring care	Sea anemones, cubozoan jellyfish, parasitoid wasps, saw flies	[55,61–63]
Mating	Scorpions	[27]
Habitat creation	Ants	[64]
Antimicrobial ointment	Ants, wasps	[65]
Ectoparasite deterrent	Slow loris	[66]
Antivenom	Tawny crazy ant (Nylanderia fulva)	[28]
Prey homing device	Rattlesnakes	[67]
Intraspecific communication	Ants, wasps	[68–70]

# How widespread is venom?

- Evolved independently more than 100 times
- Eight separate phyla

(Schendel et al., 2019)

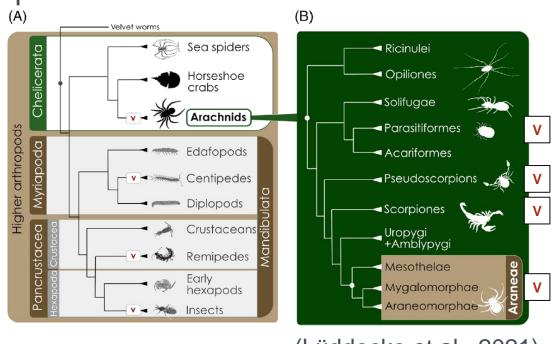


# How widespread is venom?

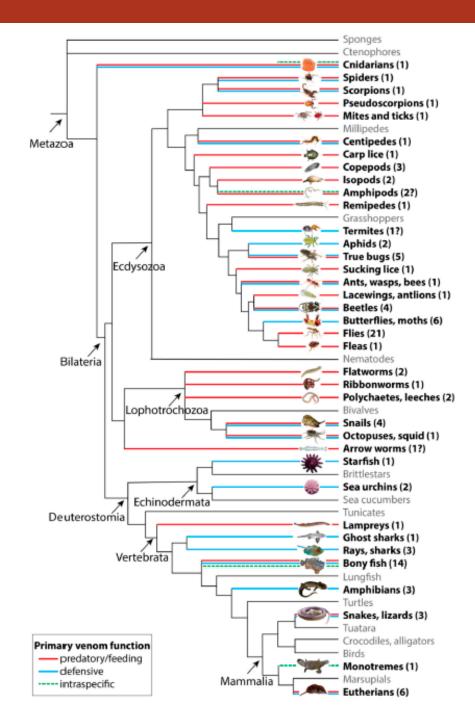
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(Schendel et al., 2019)

• Arthropods and arachnids:



(Lüddecke et al., 2021)

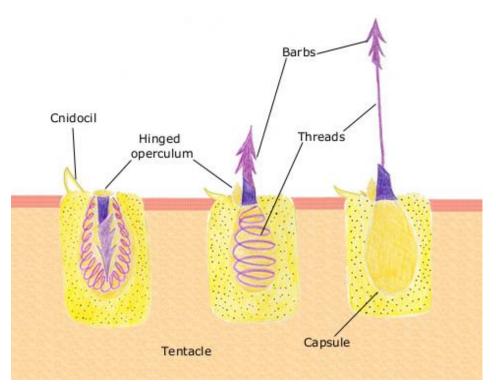


# Part 2: Venom apparatus morphology

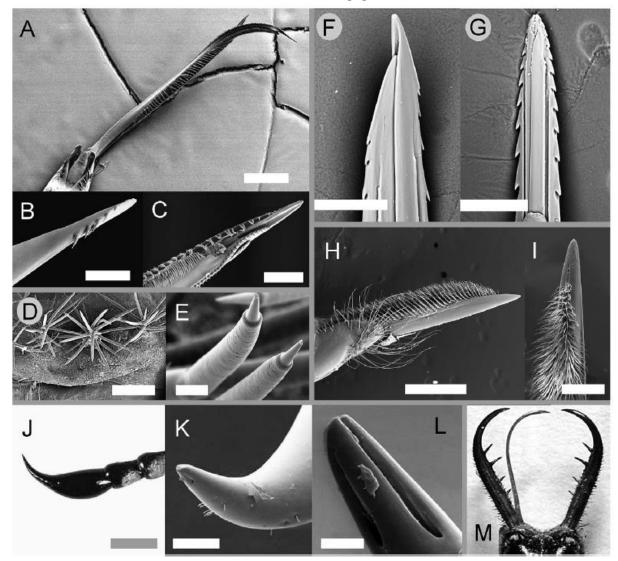


## **Gereral morphology**

- Venom glands or organelles
- Venom apparatus



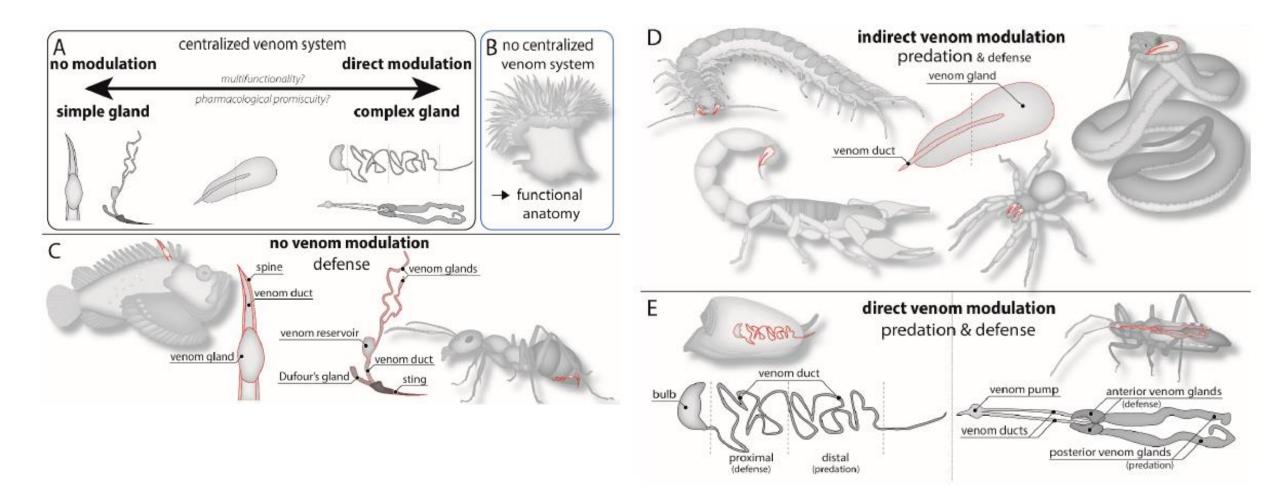
**Insect venom apparatus** 



Nematocyst of cnidarians

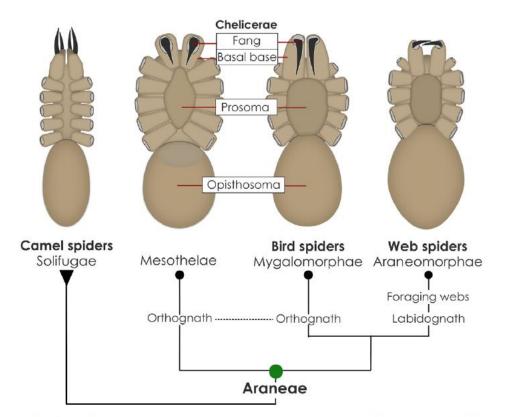
Walker al., 2018

### **Gereral morphology**



Schendel et al., 2019

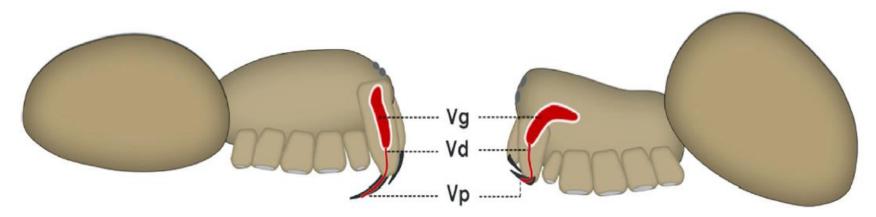
### Spider venom apparatus morphology



**Fig 2.** The chelicerae of the three major lineages within Araneae. The abstract ventral sides illustrate the different chelicerae configurations and major traits are plotted on the phylogram. Solifugae are added as an older chelicerate group that still feature the more ancient (plesiomorphic) state of chelicerae, which were originally three-segmented and scissor-like.

Lüddecke et al., 2021

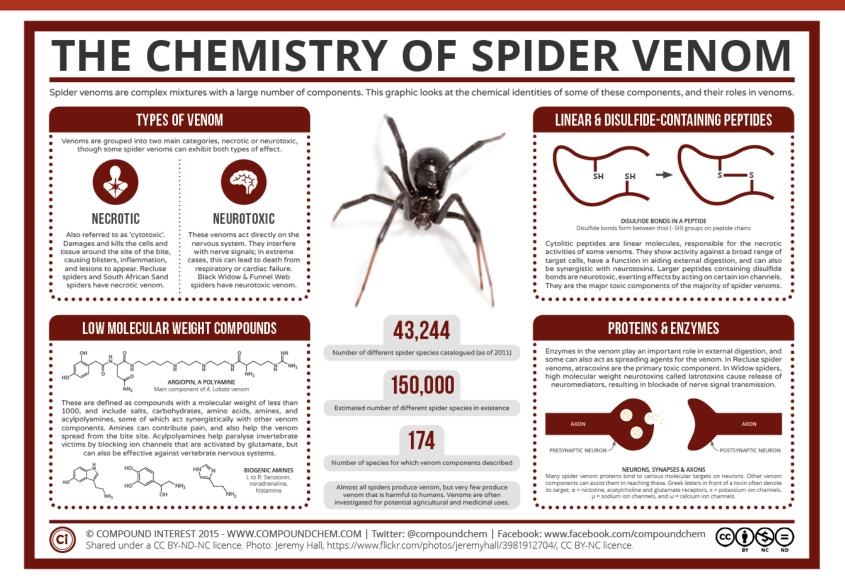
### Spider venom apparatus morphology



**Fig 3.** Comparison of the venom apparatus in orthognath and labidognath spiders. Venom glands (Vg) and venom duct (Vd) that leads to the venom pore (Vp) at the outer side of the fang tip are shown in red. The muscle layer surrounding the gland is illustrated in white.

Lüddecke et al., 2021

# Part 3: Venom chemistry



# Type of toxins

#### Haemotoxins

- disrupt haemostatic system
- not common in spiders

### • Cytotoxins

- impair the structure and function of cell membranes
- not common in spiders

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recluse spiders (Loxosceles spp.)

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# Type of toxins

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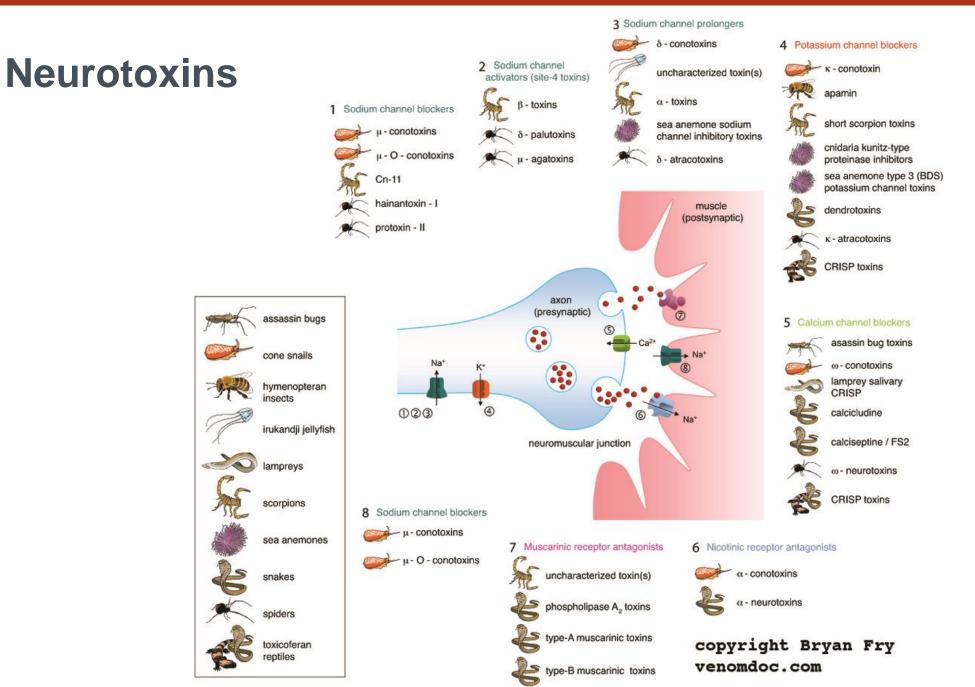
### • Cytotoxins

- impair the structure and function of cell membranes
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#### • Neurotoxins

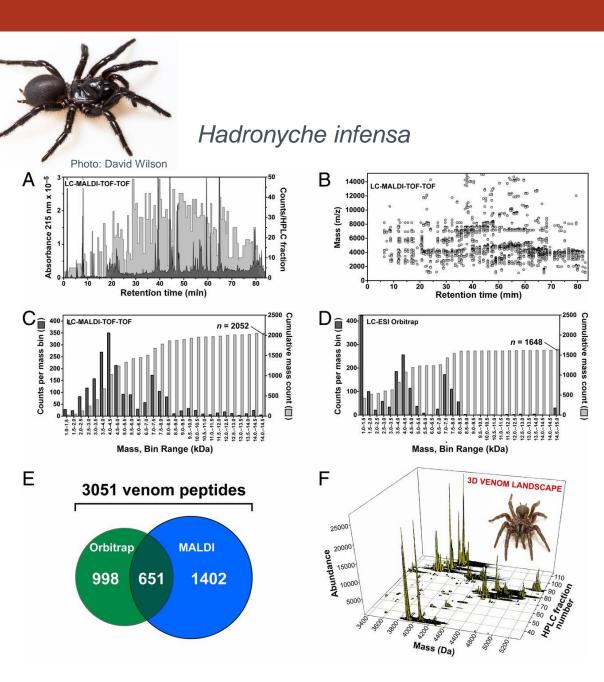
- presynaptically or postsynaptically affect neurotransmission
- main components of spider venom

#### **VENOM CHEMISTRY**



# Venom biochemistry

- Tens to thousands of components in single species
- Hereafter emphasis on spider venom
- Types of molecules:
  - Small molecules
  - Peptides
    - Antimicrobial Peptides
    - Cysteine-rich Peptides
  - Proteins



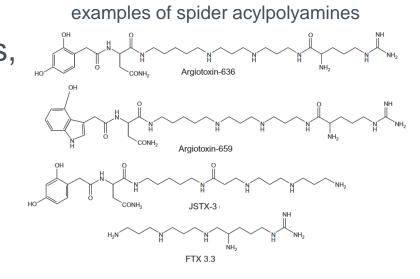
Pineda et al., 2020

# **Small molecules**

- < 1 kDa
- various compounds: ions, salts, organic acids, nucleotides, nucleosides, amino acids, amines, alkaloids and polyamines

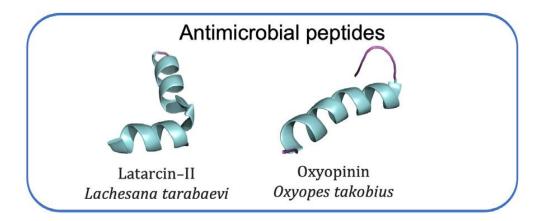
• various functions:

- neurotransmitters
- co-factors facilitating the folding and activity of toxins
- insecticidal neurotoxins



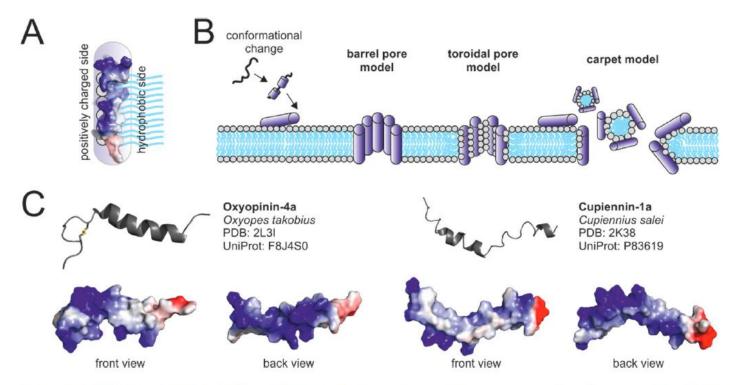
# **Peptides**

- < 10 kDa
- Antimicrobial Peptides
  - linear, α-helical peptides without disulfide bonds
  - dual function:
    antimicrobial activity or lytic peptides
  - most of those identified AMPs found in Lycosidae and Theraphosidae; also Zodariidae or Oxyopidae
  - more than 50 such peptides in the venom of Lycosa sinensis



# Peptides

- < 10 kDa
- Antimicrobial Peptides



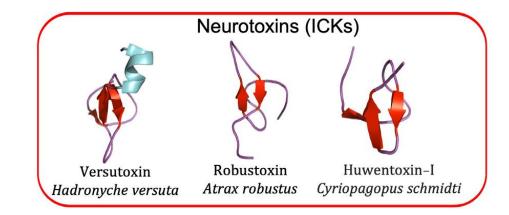
**Figure 1.** Antimicrobial peptides (AMPs) and their proposed mechanism of action. (**A**) Model of interaction between an AMP and phospholipids. AMPs assume an amphipathic α-helical structure in proximity to cellular membranes. The hydrophobic side of the helix (white) inserts into the membrane and interacts with the phospholipid side chains. The positively charged side (blue) interacts with negatively charged lipid head groups. (**B**) Models of membranolytic actions of AMPs. (**C**) NMR-based 3D structures of two antimicrobial peptides from spider venom. Electrostatics were computed using PDB2PQR [70]. Blue surfaces represent positively charged surfaces; red negative charged; and white neutral.

#### Langenegger, Nentwig & Kuhn-Nentwig, 2019

# **Peptides**

- < 10 kDa
- Antimicrobial Peptides
- Cysteine-rich Peptides
  - functionally most important group of components
  - principal neurotoxic components
  - typically with molecular masses below 10 kDa
  - rich in disulfide bonds
  - different families:

Kunitz peptides, HAND peptides, DDH peptides, ICK peptides



#### **VENOM CHEMISTRY**

# **Peptides**

- ICK (inhibitor cysteine knot) peptides
  - structure:

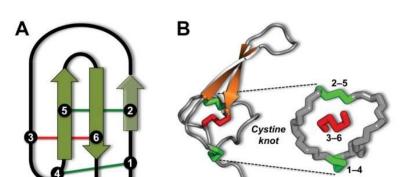
(triple-stranded) antiparallel  $\beta$ -sheets

at least 6 cysteine residues (forming 3 disulfide bridges)

- -> pseudoknot motif
- expanded cysteine scaffolds and/or double ICK (dICK) motifs
- exceptional stability
- mode of action: stable complexes with prey receptors, disrupting their normal function

#### • targets:

voltage-gated sodium, potassium and calcium channels acid-sensing ion channels, glutamate receptors transient receptor potential channels



Inhibitor cystine knot (ICK)

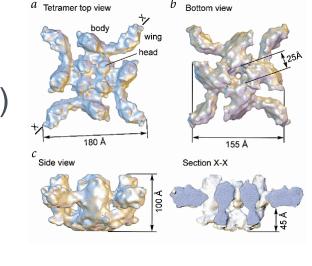
Herzig & King, 2015

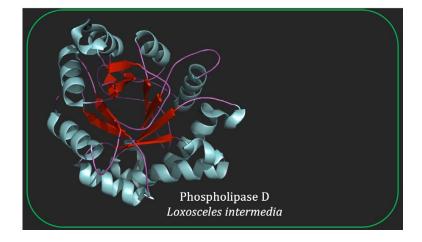
# **Proteins**

- > 10 kDa
- key venom components in some taxa (e.g., black widows)

#### latrotoxins

- homotetrameric pores in the presynaptic neuronal membranes
- phospholipase D
  - highly cytotoxic sphingomyelin-hydrolysing enzyme
- neprilysin metalloproteases, CAP proteins
  - unclear function

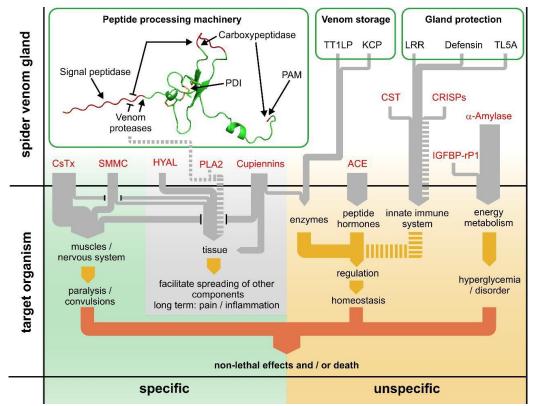




#### α-latrotoxin

# Mode of action - synergistic effects of venom compounds

- temporally and spatially regulated interactions
- dual prey inactivation strategy (Kuhn-Nentwig et al., 2019)



reviewed in Lüddecke et al., 2021

## Mode of action - synergistic effects of venom compounds

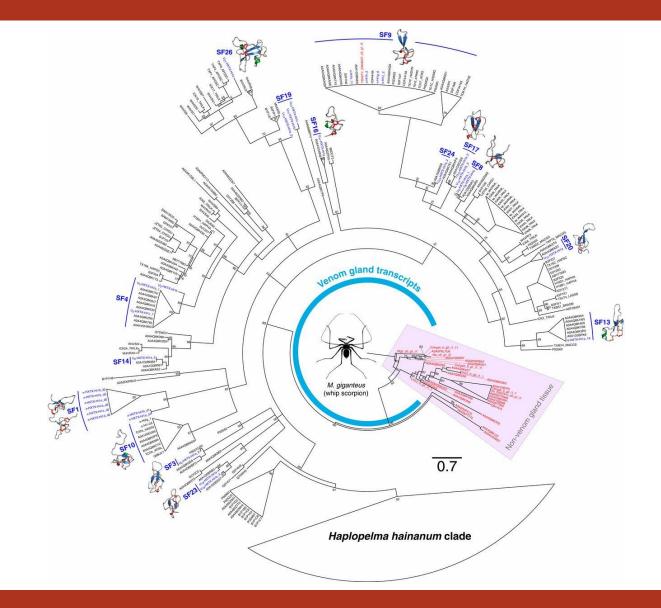
- temporally and spatially regulated interactions
- dual prey inactivation strategy (Kuhn-Nentwig et al., 2019)

Some of the large proteins immediately disrupt prey physiology and metabolism, while others act to spread the neurotoxins and thus trigger a subsequent wave of paralysis.

- some peptides mediate or enhance the bioactivity of others
- rapid paralysis followed by long-term immobilization
- neurotransmitters making binding sites accessible for other toxins

reviewed in Lüddecke et al., 2021

# Part 4: Venom evolution



Pineda et al., 2020

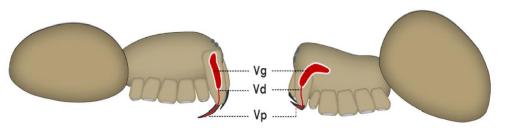
## **Evolution of the delivery system in spiders**

#### • origin not clear

- salivary glands, similar to those of ticks
- silk-producing glands present in early chelicerates

#### the size and placement of the venom apparatus

- modern araneomorphs vs Mesothelae and Mygalomorphae
- migration of the venom glands from the basal part of the chelicerae into the prosoma
- reflected during ontogenesis



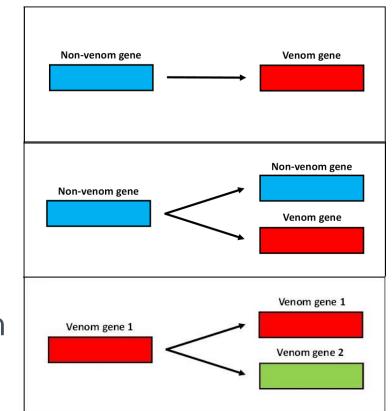
**Fig 3.** Comparison of the venom apparatus in orthognath and labidognath spiders. Venom glands (Vg) and venom duct (Vd) that leads to the venom pore (Vp) at the outer side of the fang tip are shown in red. The muscle layer surrounding the gland is illustrated in white.

reviewed in Lüddecke et al., 2021

# **Recruitment and neofunctionalization**

### **Evolution of venom compounds**

- recruitment and weaponization
  - signalling molecules into unregulated agonists or inhibitors
- frequent duplications
- existing toxins can also undergo neofunctionalization
  - new activities and functions



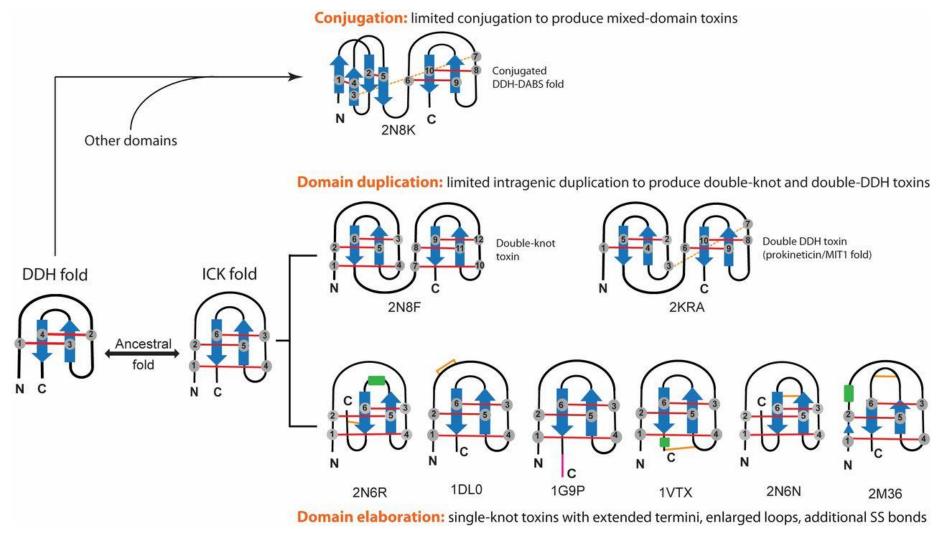
reviewed in Casewell et al., 2012; Lüddecke et al., 2021

# **Gene duplication**

#### ICK peptides

- pseudoknot motif
  - amino acid substitutions can accumulate with little impact on structure
- descendants of a single weaponized ICK lineage
- duplication and structural diversification
- domain duplication dICK peptides
- also proposed for latrotoxins

### **Overview of spider ICK peptides evolution**

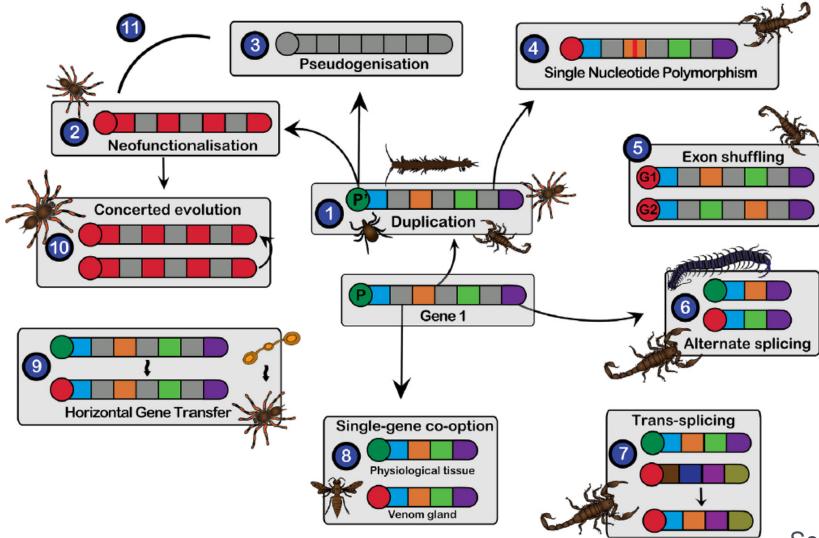


## Horizontal gene transfer

- recruitment of toxins from bacterial and fungal donors
- phospholipase D
  - in the family Sicariidae
  - a single proteobacterial ancestor
- also proposed to explain the origin of αLTX of *Parasteatoda tepidariorum*
- otherwise documented in other venomous taxa, such as centipedes (Undheim & Jenner, 2021)

reviewed in Lüddecke et al., 2021

#### **Molecular mechanisms of venom evolution - overview**



Senji Laxme et al. 2019

### Selection pressures acting on venom

- strong positive selection
- 'Red Queen hypothesis' predator *vs* prey
- but older lineages such as spiders show signatures of purifying selection
- a two-speed model of venom gene evolution
  - positive selection mostly acts during the early stages of ecological specialization
  - followed by an extended stage of purifying selection



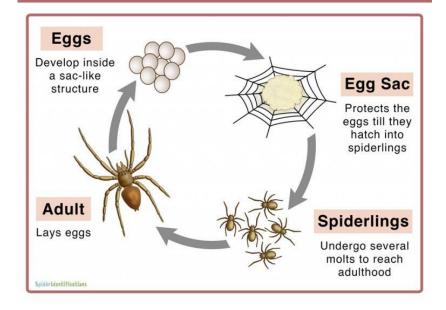
# Part 5: Venom Ecology



# Ontogeny

- venom properties linked to life-history stages
- venom yield increases as the spider ages
- venom yield declines prior to ecdysis
- compositional alterations as the spider ages

#### Spider Life Cycle



# **Geograpic variation**

- the variability of venom profiles between allopatric populations of the same species
- well documented in other venomous taxa, such as snakes
- little attention in spiders (contrasting results)
- Eratigena agrestis (Agelenidae)
  - Europe vs North America
  - no differences
- Loxosceles rufescens (Sicariidae) Latrodectus spp. (Theridiidae)
  - differences in venom profiles and toxicity

Eratigena agrestis



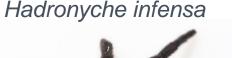
Photo: Rudolf Macek

Loxosceles rufescens



# **Sexual dimorphism**

- well documented in spiders
- australian funnel-web spiders
  - fatal males bites in humans, females much less toxic
- Phoneutria nigriventer (Ctenidae)
  Loxosceles intermedia (Sicariidae)
  - females more toxic than males
  - sex-specific components





Phoneutria nigriventer



Photo: Graham Wise

# Individual variability

- well documented in other venomous taxa, less attention in spiders
- only recently documented in Hadronyche valida



### **Envenomation strategies**

- envenomation as major strategy to incapacitate prey in most spiders
  - some exceptions: Araneidae, Uloboridae, Scytotidae
- toxins on the silk strands of the web (Esteves et al., 2020)
- venom in defence
  - escape prioritized
  - dry bites
  - aposematism

(supported by pain-inducing components)

Uloborus walckenaerius



Argiope bruennichi



Photo: Christine Hanrahan

Scytodes thoracica



Photo: Fritz Geller-Grimm

# **Venom optimization**

- venom physiologically expensive
- venom optimisation (or venom metering)
  - conservation of venom resources
  - economical delivery of venom
    - weak prey -> small amount of venom
    - resistant/dangerous prey -> larger ammount of venom

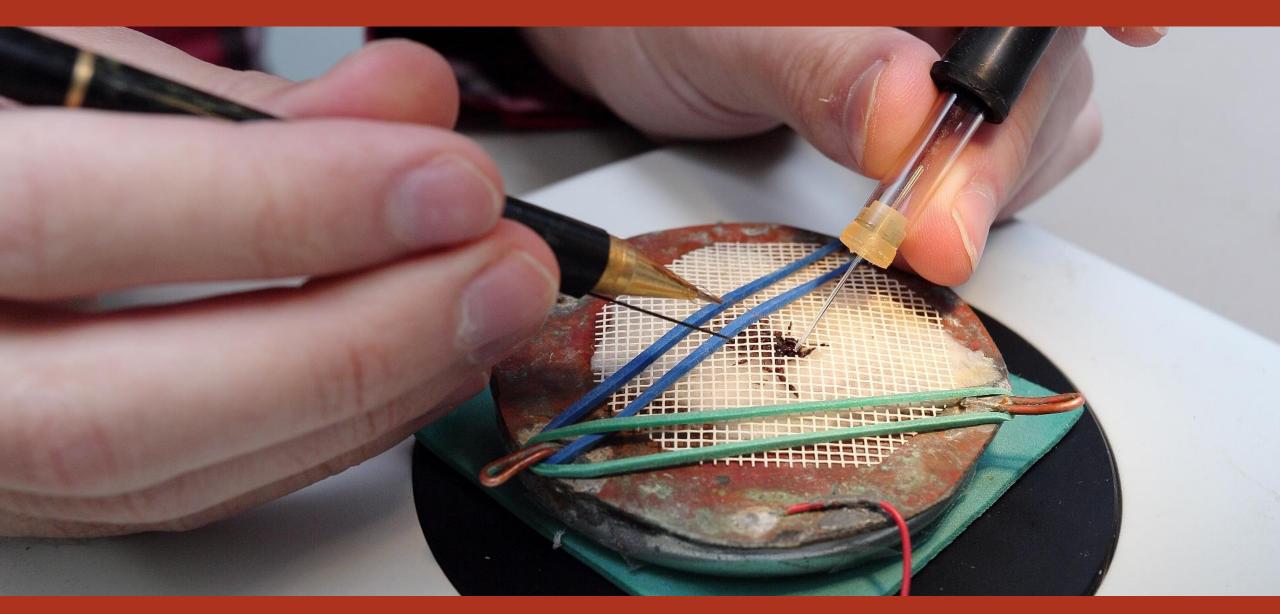
#### trophic specialisation

- highly effective, but simpler venom
- dispensable venom components purged from the venom to save resources

Lüddecke et al., 2021; Morgenstern & King 2013; Pekár et al., 2018a,b



# Part 6: Methods & Applications



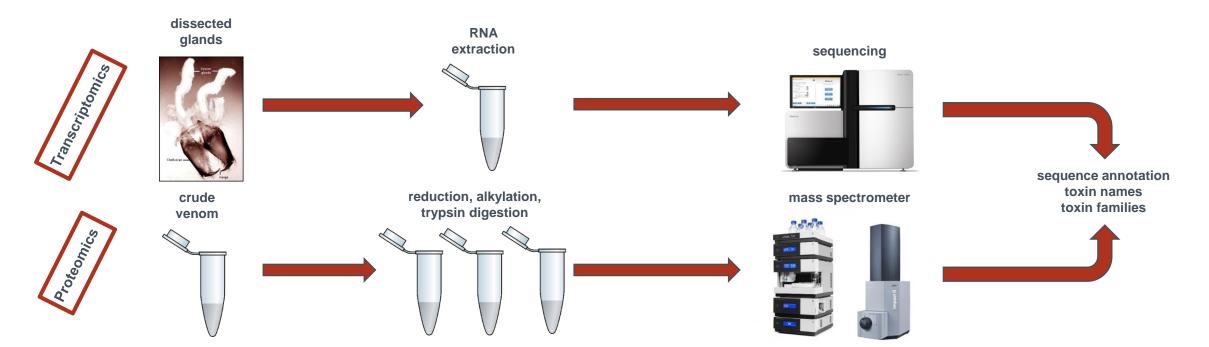
### **Behavioural experiments**

- laboratory experiments with living specimens
- for example, observation of prey paralysis after bite (e.g., Pekár et al, 2018b)



### **Venom composition - venomics**

- integration of transcriptomic, proteomic (and genomic) approaches
- transcriptomics of the venom-producing tissue
- mass spectrometry (MS) based proteomics
- bioinformatic integration of the data



# Structure of venom compotents

- X-ray crystallography
  - rarely used for spider toxins
  - larger proteins
- nuclear magnetic resonance (NMR)
  - most used for spider toxins
  - peptides
- cryoelectron microscopy (EM)
  revelue used for epider toxing
  - rarely used for spider toxins



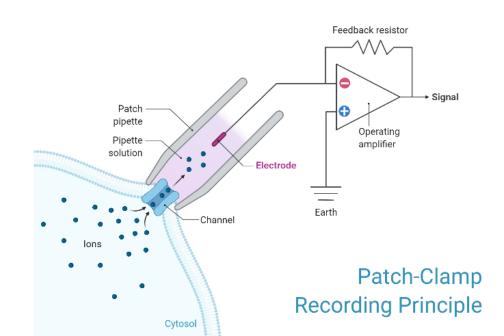
# Venom physiology bioassays

#### venom efficiency

- venom injection bioassays
- crude venom, venom fractions, isolated recombinant toxins

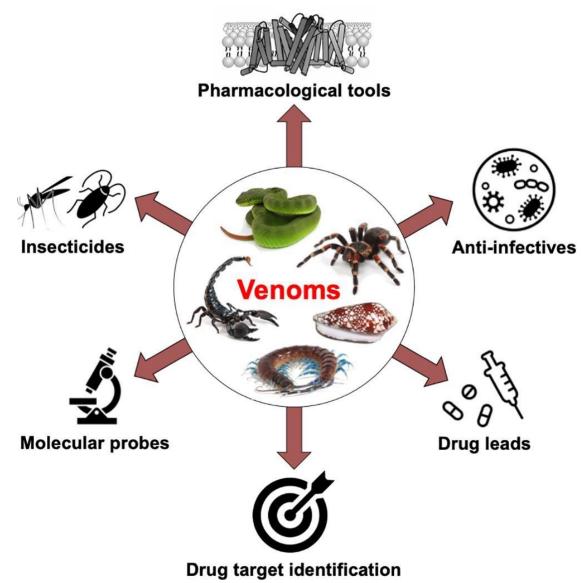
#### venom physiology

- patch clamp technique
- target Ion Channels synthesized and expressed in *Xenopus* oocytes
- toxins added (recombinant toxins)
- electrophysiological two-electrode voltage-clamp recordings



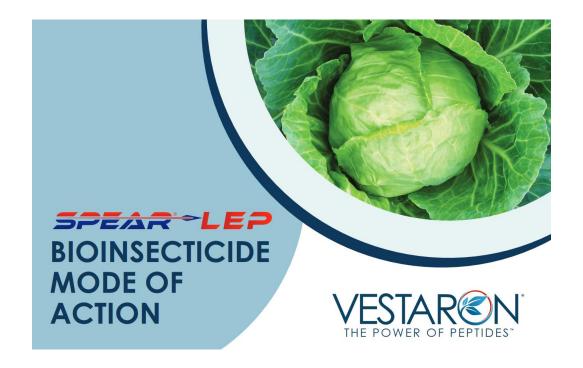


## **Venom applications**



### **Venom applications - pesticides**

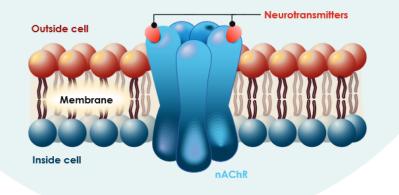
eco-friendly bioinsecticides

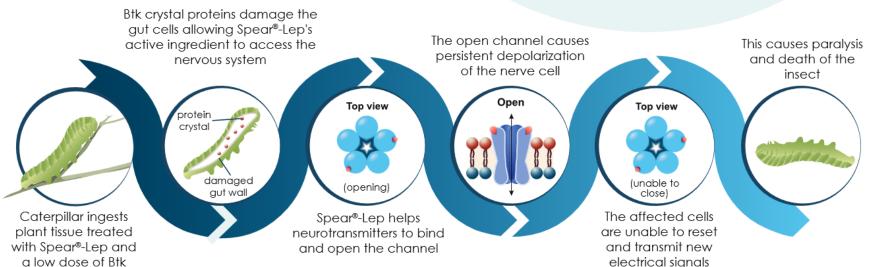


#### **Venom applications - pesticides**

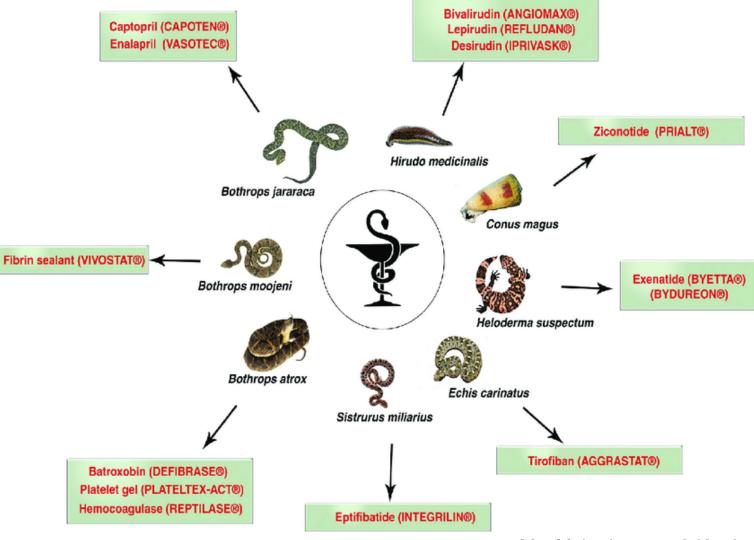
The active ingredient in Spear® is GS-omega/ kappa-Hxtx-Hv1a. Spear® products are the first peptide-based insecticides, and the first bioinsecticides that affect a specific neuromuscular target. Spear® delivers an entirely new mode of action for crop protection (IRAC group 32), which means no cross resistance to any other active ingredient, and a novel tool for insecticide resistance management. Because of its biological origins, Spear® is lethal to insect and mite pests, but non-toxic to bees, fish and mammals.

Nicotinic acetylcholine receptors (nAChR) are channels found in the nerves of insects that respond to neurotransmitters. These receptors are essential for transducing certain electrical signals, such as the muscle contraction.





#### **Venom applications - drug leads**



Ma, Mahadevappa & Kwok, 2015

### **Venom applications - drug leads**



prof. Glenn King



funnel-web spider Hadronyche infensa

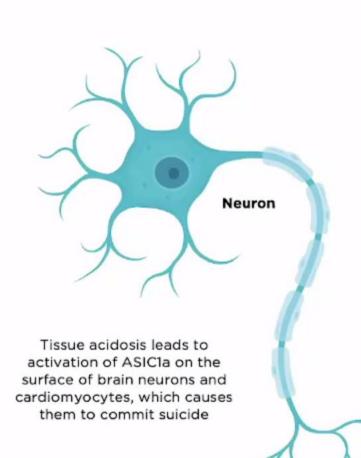
THE UNIVERSITY OF QUEENSLAND AUSTRALIA Institute for Molecular Bioscience

-> cure for stroke

#### ASIC1a MEDIATES CELL DEATH AFTER STROKE & MI

Cardiomyocyte

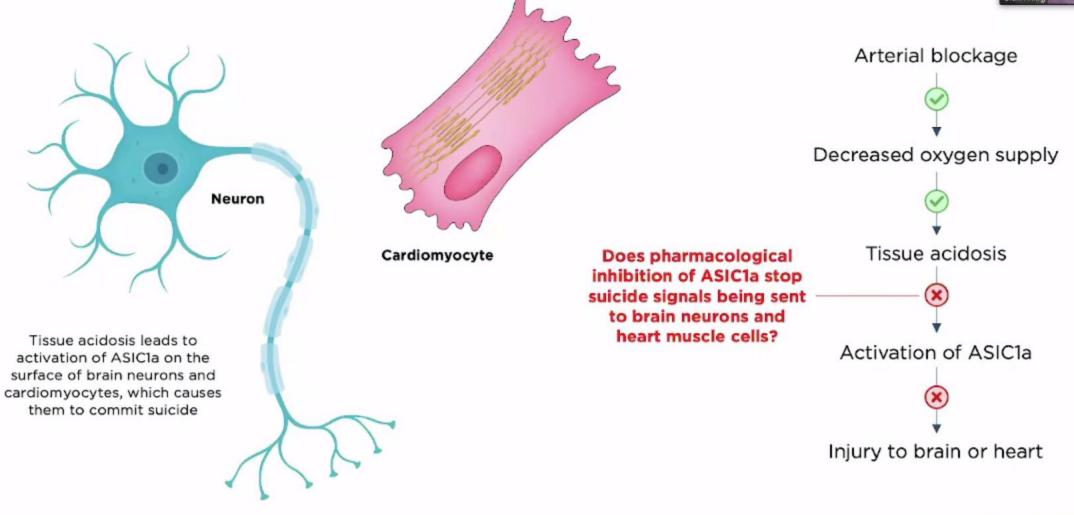






#### ASIC1a MEDIATES CELL DEATH AFTER STROKE & MI

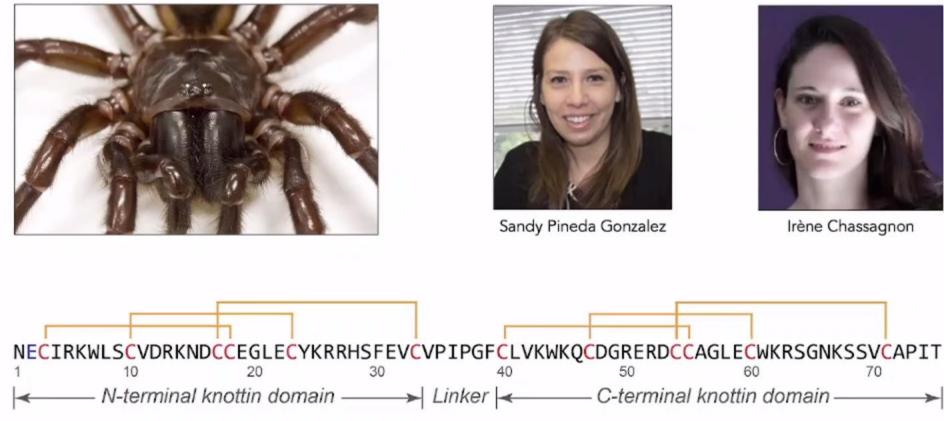




**1ST EUVEN INTERNATIONAL CONGRESS** 

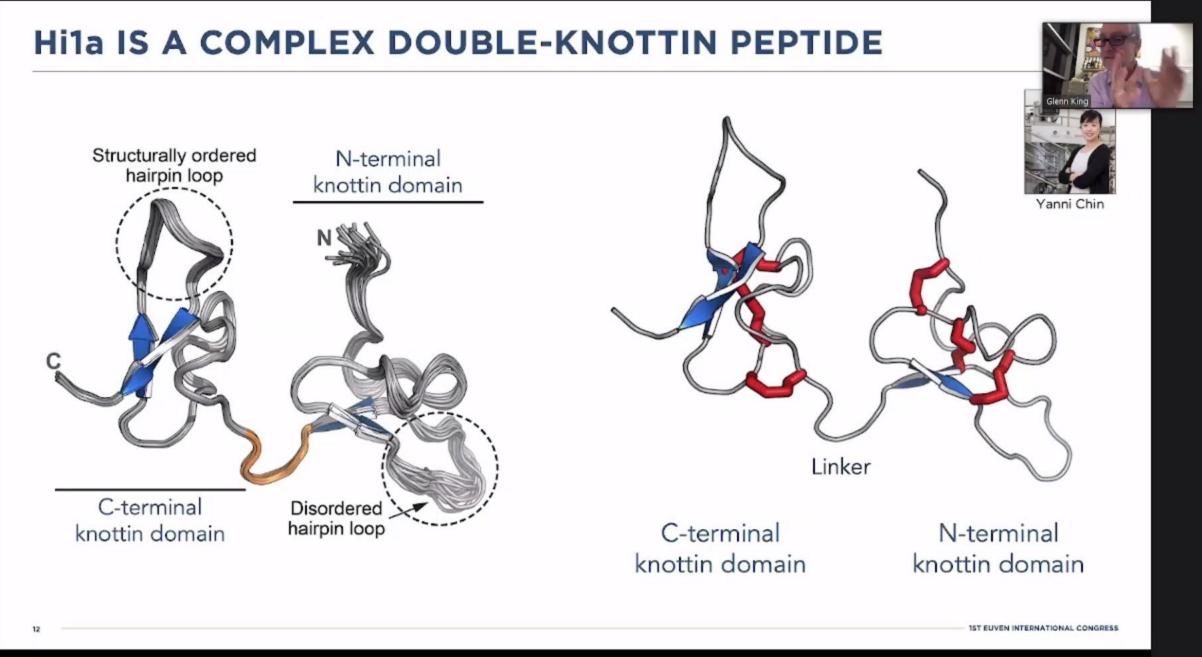
#### **SERENDIPITOUS DISCOVERY OF Hi1a**



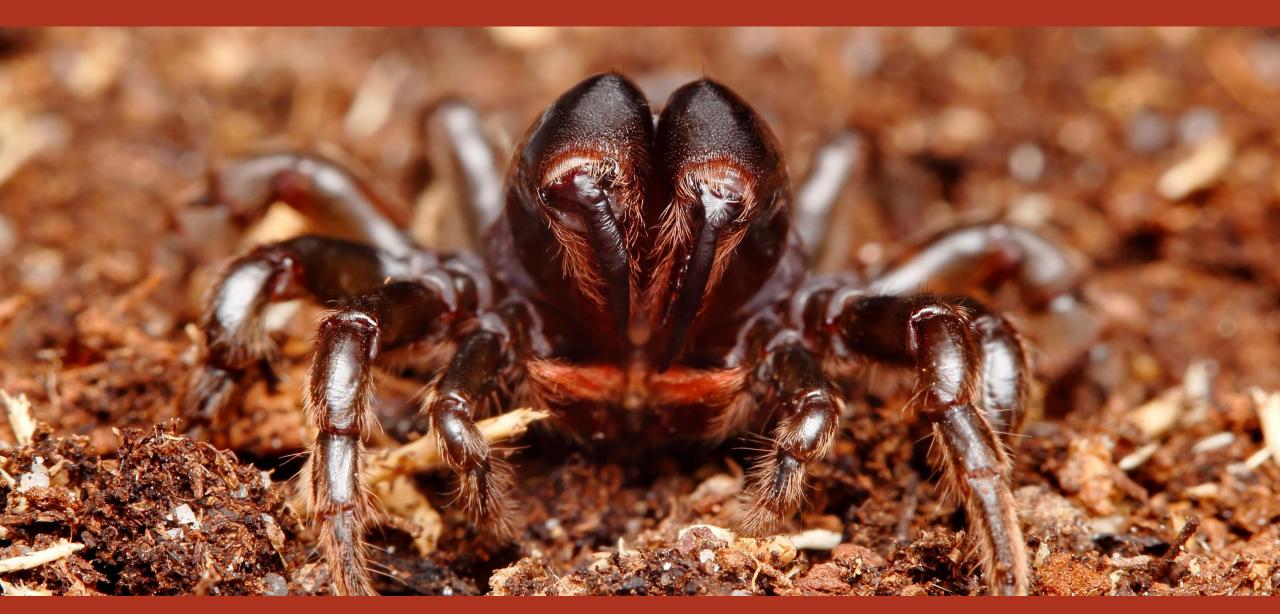


Hi1a: 75 residues; 6 disulfide bonds = 10,395 disulfide isomers

1ST EUVEN INTERNATIONAL CONGRESS



# **Summarization**



# **Summarization**

- one of the key traits of spiders
- mainly for predation and defence
- hundreds to thousand compounds per species
- neurotoxins, mainly ICK peptides
- evolved through duplication and neofunctionalization
- positive selection followed by purifying selection
- sexual dimorphism, perhaps some geographic and individual variability
- costly substance venom optimisation
- various methods to study venoms; promising applications (drugs and pesticides)

# REFERENCES

Arbuckle, K. (2017). Evolutionary context of venom in animals. *Evolution of venomous animals and their toxins*, 24, 3-31.

Casewell, N. R., Wüster, W., Vonk, F. J., Harrison, R. A., & Fry, B. G. (2013). Complex cocktails: the evolutionary novelty of venoms. *Trends in ecology & evolution*, *28*(4), 219-229.

Esteves, F. G., dos Santos-Pinto, J. R. A., Ferro, M., Sialana, F. J., Smidak, R., Rares, L. C., ... & Palma, M. S. (2020). Revealing the Venomous Secrets of the Spider's Web. *Journal of Proteome Research*, *19*(8), 3044-3059.

Langenegger, N., Nentwig, W., & Kuhn-Nentwig, L. (2019). Spider venom: components, modes of action, and novel strategies in transcriptomic and proteomic analyses. *Toxins*, *11*(10), 611.

Lüddecke, T., Herzig, V., Von Reumont, B. M., & Vilcinskas, A. (2021). The biology and evolution of spider venoms. *Biological Reviews*.

Malaque, C. M. S., Chaim, O. M., Entres, M., & Barbaro, K. C. (2016). Loxosceles and loxoscelism: biology, venom, envenomation, and treatment. *Spider venoms. Dordrecht: Springer Nature*, 419-44.

Morgenstern, D., & King, G. F. (2013). The venom optimization hypothesis revisited. *Toxicon*, 63, 120-128.

Nelsen, D. R., Nisani, Z., Cooper, A. M., Fox, G. A., Gren, E. C., Corbit, A. G., & Hayes, W. K. (2014). Poisons, toxungens, and venoms: redefining and classifying toxic biological secretions and the organisms that employ them. *Biological Reviews*, *89*(2), 450-465.

Pekár, S., Bočánek, O., Michálek, O., Petráková, L., Haddad, C. R., Šedo, O., & Zdráhal, Z. (2018a). Venom gland size and venom complexity—essential trophic adaptations of venomous predators: a case study using spiders. *Molecular ecology*, *27*(21), 4257-4269.

Pekár, S., Líznarová, E., Bočánek, O., & Zdráhal, Z. (2018b). Venom of prey-specialized spiders is more toxic to their preferred prey: A result of prey-specific toxins. *Journal of Animal Ecology*, *87*(6), 1639-1652.

Pineda, S. S., Chin, Y. K. Y., Undheim, E. A., Senff, S., Mobli, M., Dauly, C., ... & King, G. F. (2020). Structural venomics reveals evolution of a complex venom by duplication and diversification of an ancient peptide-encoding gene. *Proceedings of the National Academy of Sciences*, *117*(21), 11399-11408.

Senji Laxme, R. S., Suranse, V., & Sunagar, K. (2019). Arthropod venoms: Biochemistry, ecology and evolution. *Toxicon*, *158*, 84-103.

Schendel, V., Rash, L. D., Jenner, R. A., & Undheim, E. A. (2019). The diversity of venom: the importance of behavior and venom system morphology in understanding its ecology and evolution. *Toxins*, *11*(11), 666.

Walker, A.A., Robinson, S.D., Yeates, D.K., Jin, J., Baumann, K., Dobson, J., Fry, B.G. and King, G.F. (2018). Entomo-venomics: The evolution, biology and biochemistry of insect venoms. *Toxicon*, 154, 15-27.