

## Anti-Bacterial Effect of Venom Extracted from *Pardosa oakleyi* (Family Lycosidae) and Silk Retrieved from *Crossopriza lyoni* (Family Pholcidae)

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Received: June 05, 2019; Published: September 16, 2019

DOI: 10.31080/ASMI.2019.02.0373

### Abstract

Spiders are widely known and abundantly present successful predators because they are surprisingly armed with potent cocktail of venom along with the multifunctional tangled webs. In current research, venom was recovered from *Pardosa oakleyi* (Family Lycosidae) and silk was recovered from *Crossopriza lyoni* (Family Pholcidae). Study was aimed to partially characterize the venom of *P. oakleyi* and for estimation of antibacterial potency of venom and silk. Four pathogenic strains of bacteria were used i.e., Gram positive (*Staphylococcus* sp. and *Streptococcus* sp.) and Gram negative (*Acinetobacter* sp. and *Pasteurella* sp.). Results revealed that the venom of *P. oakleyi* comprised of relatively high molecular weight peptides ranging from 155kDa to 43kDa. And susceptibility tests indicated that the crude venom was ineffective against all tested strains. Although the silk of *C. lyoni* exhibited significant bacteriostatic action against *Acinetobacter* sp. and *Streptococcus* sp.

**Keywords:** Characterization; Bacteriostatic Action; Peptides; Susceptibility Test; Crude Venom; Gram Positive and Gram Negative Strains

### Abbreviation

SDS PAGE: Sodium Dodecyl Sulfate Polyacrylamide Gel Electrophoresis.

### Introduction

The rapid development of resistance in bacteria is taking place throughout the world which is threatening the effectiveness of antibiotics, the savior of millions of lives [1,2]. Several decades after initially patients were cured using antibiotics, bacterial infections have again turned into a tenacious and alarming threat [3]. Excessive utilization, mishandling and unsystematic use of broad-spectrum antibiotics are the major contributors in the emergence of resistivity among bacterial strains [4,5]. Almost all clinically significant bacterial pathogens show resistivity against antibiotics. The decrease in effectiveness of antibiotics in treating common infec-

tions and the advent of novel untreatable strains indicates that we are at the edge of a post-antibiotic era [6].

The regular and prompt development of new interventions with antibacterial activity is the need of time to deal all challenges related to antibiotic resistivity in bacteria. Natural products are surpassing the synthetic products and providing foundation for innovation of chemical diversities including natural antibiotics [7]. These natural antibiotics are reported to be more effective, safer and have less side-effects on human health and environment as compared to synthetic antibiotics [8].

In phylum Arthropoda, spiders are one of the most diverse group with approximately 100,000 existing species [9]. Spiders

are amazingly armed with potent cocktail of venom along with the multifunctional tangled webs. Spider venom consist of a wide range of biologically active compounds including various peptides, proteins, enzymes, salts and small organic molecules [10,11]. All the toxins in spider venom are intended to block variety of receptors, channels, membranes and diverse enzymes of invertebrates and vertebrates [12].

In past minor work has been done on isolation and applications of peptides extracted from spider venom. Owing to this insufficiency in research, the number of discovered peptides is reasonably less [11]. Usually spider venom contains peptides up to 25% by weight with great diversity of almost 12 million types of discovered peptides [13,14]. Moreover, peptidic content of venom is highly selective or specific in targeting. This property of venom peptides can lead to development of novel therapeutics [15,16].

Moreover, spider silk comprises off diverse compounds such as potassium nitrate, bisphosphonate peptides [17]. Presently, spider silk proof itself a marvelous gift of nature by exhibiting its multiple potentials. It is wonderful combination of strength and extensibility [18]. Spider silk can be amazingly useful in medical field because of its compatibility with living tissues of humans [17,18] and most importantly spider silk contain number of antimicrobial compounds such as potassium nitrate, bisphosphonate peptides and phospholipids hydrates with obvious antibacterial potential [19,20]. Another important constituent of spider silk is potassium hydrogen phosphate. It basically releases protons in aqueous medium to make the silk acidic (pH 4), which inhibit the microbial growth [21]. Furthermore, the lipid present in spider silk holds 12-methyltetradecanoic acid and 14-methylhexadecanoic acid that restrict microbial growth [21]. Silk contains active agents which do not allow bacterial strains to gain resistivity against natural antibacterial agent [22]. These properties make spider silk an admirable candidate for innovation in antimicrobial drug discovery.

Hence the venom and silk, extracted from spiders, could be one of the most effective ones among fewer natural resources to be discovered as antimicrobial drugs, as they are surprisingly remarkable weapons of spiders and so far very little or no attention has been paid to them in this context.

Therefore, this study was conducted in order to check antibacterial potential of potent peptides present in spider venom (*Pardosa oakleyi*) and spider silk (*Crossopriza lyoni*). For this purpose partial characterization of venom (*Pardosa oakleyi*) was also done and to evaluate antibacterial potential of venom and silk four different pathogenic strains (including gram positive and gram negative both) were used. Significant bacteriostatic property by these peptides may lead to invention of novel therapeutics.

## Materials and Methods

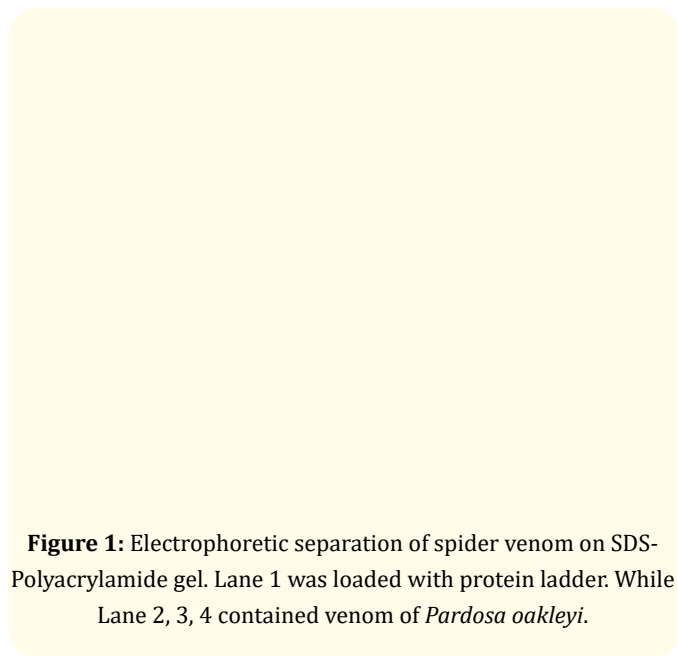
- **Spider collection and venom extraction:** Spiders (*Pardosa oakleyi*) were collected from citrus orchard and agricultural fields. Spiders were collected in separate aerated vials to avoid cannibalism among them. Chelicerae along with venom glands were separated after immobilization of spiders through freezing [23]. After that crude sample was prepared by homogenizing extracted chelicerae in 0.5ml of tris HCl buffer of pH 8.2 and molarity 0.05 M [24] and was centrifuged (MPW-352R) at 15,000 rpm for 20 minutes at 4°C. Then supernatant extracted through suction pipette was frozen at -20°C in Ultra freezer [24].
- **Characterization of venom by sodium dodecyl sulphate (SDS) gel electrophoresis:** Characterization involves fractionation of a variety of constituent proteins on the basis of their molecular weights. SDS polyacrylamide gel electrophoresis was performed to separate different venom fractions by following protocol of Sambrook and Russle [25]. Gel was categorized as 5% stacking gels and 12% resolving gel on the basis of different percentages. Supernatant containing venom sample melted within 5 minutes and were mixed in equal ratio (1:1) with 2X SDS loading buffer (100M tris-Cl, pH 6.8) before heating at 100°C.
- After running the sample with ladder (EZ™-Prestained protein Ladder Marker), gel was stained in Comassi brilliant blue staining solution for 1 hour and then shifted to destaining solution (40% methanol and 30% acetic acid) for 6-8 hours. Stained gels were photographed and approximate molecular weight of different peptide fractions were revealed by comparing with reference protein standards.
- **Silk collection and silk solution preparation:** Approximately 30 mg of silk was extracted from the selected spider species, *Crossopriza lyoni*. To prepare stock silk solution, 30 mg of silk was dissolved in 5% NaOH by heating

- **Susceptibility test:** For the examination of antibacterial potency of spider venom extracted from *Pardosa oakleyi* and spider silk recovered from *Crossopriza lyoni*, a susceptibility test following Kirby-Bauer Disk Diffusion method was used. Obvious inhibition zones were appeared against two Gram negative (i.e *Acinetobacter sp.* and *Pasteurella sp.*) and two are Gram positive (i.e *Staphylococcus sp.* and *Streptococcus sp.*) strains. These strains were cultured on nutrient Agar and maintained in liquid nutrient broth at 37°C.
- **Statistical analysis:** The Mean and the Standard error of mean (Mean  $\pm$  SE) were computed using Minitab 14. One way ANOVA followed by Tukey's test was applied to compare the zone of inhibitions of treatments using SPSS version 13 (Statistical Package for Social Sciences, 13).

## Results and Discussion

### Partial characterization of venom and determination of peptide fractions

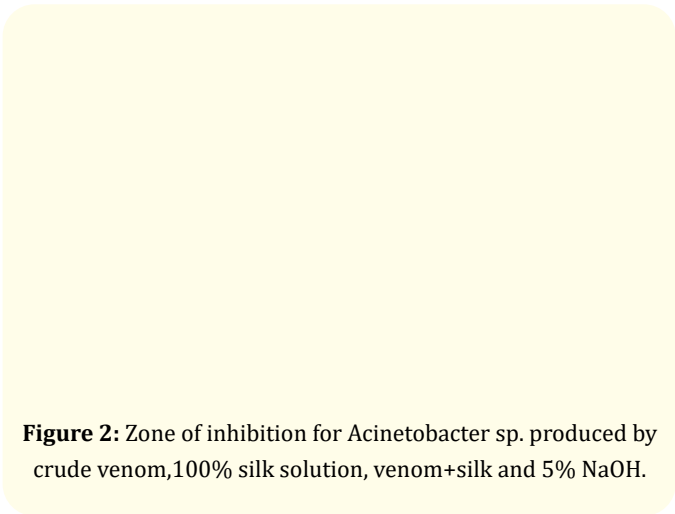
During gel electrophoresis we compared resolved protein bands to protein bands of known molecular weight, alienated on same gel. When venom of *P. oakleyi* was analyzed through gel electrophoresis, five different protein bands of 155kDa, 140kDa, 100Da, 65kDa and 43kDa were appeared. Band of 43KDa was most intense and broad among all bands followed by 100kDa, 70kDa and 155KDa. Whereas band of 140kDa was less intense (Figure 1).



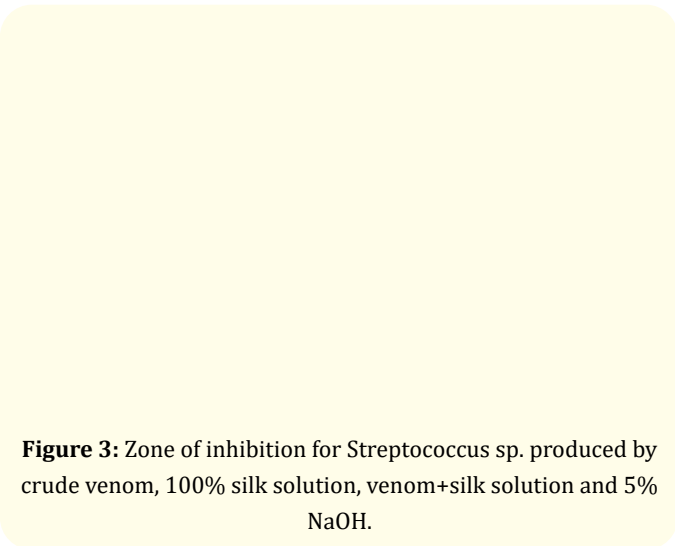
**Figure 1:** Electrophoretic separation of spider venom on SDS-Polyacrylamide gel. Lane 1 was loaded with protein ladder. While Lane 2, 3, 4 contained venom of *Pardosa oakleyi*.

### Antibacterial activity

The crude venom of *Pardosa oakleyi* did not exhibited bactericidal activity against any of the bacterial strain. Conversely by using silk solution, significant inhibition zones appeared in two of the tested bacterial strains i.e. *Acinetobacter sp.* and *Streptococcus sp.* The maximum zone of inhibition was obtained in case of *Acinetobacter sp.* (i.e.,  $3.523 \pm 0.433$ ) upon 100% silk treatment (Figure 2). The venom+silk solution induced reduced inhibition zones as compared to 100% silk. The inhibition zones shown by control groups were less significant as compared to both silk and venom+silk treated *Acinetobacter sp.* Similar results were obtained against *Streptococcus sp.* ( $P < 0.001$ ) (Figure 3). However, in the case of *Staphylococcus sp.* and *Pasteurella sp.*, both 100% silk and venom+silk solution failed to produce significant inhibition zone.



**Figure 2:** Zone of inhibition for *Acinetobacter sp.* produced by crude venom, 100% silk solution, venom+silk and 5% NaOH.



**Figure 3:** Zone of inhibition for *Streptococcus sp.* produced by crude venom, 100% silk solution, venom+silk solution and 5% NaOH.

## SDS-PAGE

In present research, extracted venom from *Pardosa oakleyi* give rise to six bands (43-155kDa) on gel. Band of 43kDa was most significant and broad among all, which indicate high concentration of these peptides. Whereas as band of 140kDa was less significant. Effectiveness and mode of action of venom peptides of similar molecular weight as toxin is reported in available literature. It is described that the venom peptides of more than 10kDa molecular weight acts as toxin [26]. Moreover, cyto-insectotoxins of molecular weight 8kDa were extracted from Central Asian spider [27]. Similarly, venomous peptides of molecular weight 10kDa get identified in *Selenocosmia huwena* [28]. Venomous proteins of molecular weight ranging from 1 to 40kDa were reported in the venom of *Loxosceles* spiders [29]. Furthermore, Australian funnel-web spider, *Hadronyche infensa* contains noxious peptides having masses between 1,000 to 15,000 Da. While cytolytic peptides are generally smaller (~3kDa) and the size of spider neurotoxins is fluctuated between 3–7kDa [30]. That's why the peptide fraction present in the venom of selected species doesn't act like neurotoxin or cytotoxic.

## Effect of spider venom on cell culture

Antibacterial potency of venom extracted from *Agelena labyrinthica* (Araneae: Agelenidae) is also reported [31]. It is demonstrated that Lycotoxins I and II (containing 25 and 27 amino acids respectively) isolated from the wolf spider *Hogna carolinensis* show potent lytic activity against Gram-negative bacteria (*Escherichia coli*) and yeast (*Candida glabrata*) [32]. Moreover, the peptides known as laticins extracted from venom of *Lachesana tarabaei* and spiderines of *Oxyopes takobius* venom exhibit cytolytic activity in gram-positive and gram-negative bacteria along with fungal cell i.e. yeast [33]. It is reported that the venom of *C. salei* comprised of lytic peptide having noticeable antibacterial potency against Gram positive bacteria i.e. *Staphylococcus epidermidis* and *B. subtilis* and Gram negative bacteria i.e. *E. coli*, *Pseudomonas putida* and *Paracoccus denitrificans* [34].

But still our current research reflects that crude venom of *Lycosids* (*Pardosa Hadronyche infensa*) has no bactericidal effect on any of the tested strain. These contradictory results can be explained by presence of peptides of high molecular weight, these larger peptides cannot penetrate cell membrane. So, that's why they cannot exhibit cytolytic activities. Here one thing is noteworthy that the

peptides present in venom of *Pardosa oakleyi* have higher molecular masses. These larger peptides cannot exhibit cytolytic activities as they cannot penetrate cell membranes.

## Effect of spider silk on bacterial culture

Currently we examined spider silk for its antibacterial activity. Results reveal inhibitory effect of *Crossopriza lyoni* silk against Gram negative *Acinetobacter* sp. and Gram positive *Streptococcus* sp. Many researchers have discussed the possibility that spider silk can induce growth inhibitions in microbes [21,34,35]. Antimicrobial effect of spider silk against both Gram negative and Gram positive strains has been reported also [36]. According to a report, silk of *Nephila pilipes* possess antibacterial activity against gram negative *E. coli* and *Pseudomonas aeruginosa* as well as against gram positive *Staphylococcus aureus* [37]. These findings are in accordance with our research, indicating that the spider silk is potent for both Gram negative as well as Gram positive strains.

However, some studies have revealed that the spider silk only inhibits Gram positive strains. A study has shown that the *Tegegnaria domestica* silk has inhibited the growth of the Gram positive bacterium [38]. *Bacillus subtilis* while has no effect on *E. coli*, a gram negative bacteria. Furthermore, the *Lasiodora parahybana* silk is shown to possess some antimicrobial activity against Gram positives [39]. This contradiction could be explained by the fact that the antimicrobial effect of silk of different spider species varies with the attributes, quality, type and the arrangement of amino acids in the silk fibre [38].

Our results also revealed that the spider silk did not inhibit the growth of *Staphylococcus* sp. and *Pasteurella* sp. This may be explained by the fact that our experimental set up was not sensitive enough to detect significant inhibitions against this microbe or it could be possible that like many other antimicrobial agents, our treatment has narrow spectra of activity and it generates differential inhibitions in different types of bacteria. The difference in inhibitions of bacteria could also be explained by the species phylogenetic distance.

Present study exposed that the spider silk more efficiently inhibit the growth of aerobic Gram negative *Acinetobacter* sp. The largest zones of inhibitions were noticed in *Acinetobacter* sp. pursued by

Gram positive *Streptococcus* sp. Studies have revealed less adherence of surface of spider silk to Gram negative bacteria (*P. aeruginosa* and *E. coli*) in contrast to Gram positive bacteria (*B. subtilis*) [40,41]. Web silk of *Nephila pilipes* reported as greater inhibitory substance against Gram negative strains (*Pseudomonas aeruginosa* and *Escherichia coli*) as compared to Gram positive strains (*Staphylococcus aureus*) [37].

The findings of current research suggesting that the spider silk is more potent against Gram negative strain is contradictory to the findings of Mirghani, *et al.* (2012), Roozbahani, *et al.* (2014) and Wright and Goodacre (2012) who showed that Gram positive strains are more susceptible to the inhibitions induced by spider silk. This contradiction can be again explained by the differential antimicrobial activity of spider silk recovered from different spider species and the phylogenetic distances of selected bacterial strains.

### Conflict of Interest

The authors declare that they have no conflict of interest.

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**Volume 2 Issue 10 October 2019**

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