

REVIEW ARTICLE

FRANCISELLA TULARENSIS - 100 YEARS: TULAREMIA RESEARCH IN FORMER CZECHOSLOVAKIA AND IN THE CZECH REPUBLIC

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Summary

The history of national tularemia research started in 1936 when the first outbreak was recognized in south-east Moravia. Since then in average about one hundred cases have been recorded annually. As tularemia was endemic in former Czechoslovakia, three research groups which concentrated on this disease were formed during decades. The first two groups have worked from sixties and were associated with Jiri Libich (Prague) and Darina Gurycova (Bratislava). The third group which concentrated on the research of natural foci started during late seventies in Valtice (Zdenek Hubalek). The experimental research was, and still is, mainly associated with military research, recently with the Proteomic Center (Faculty of Military Health Sciences, University of Defence) in Hradec Kralove. This center opens molecular approaches to the analysis of *Francisella tularensis* microbes on one side and the studies on mutual host-pathogen interaction on the other side. One of the significant aims of the research is searching for the new typing and diagnostic markers of *Francisella tularensis* for the military and medical practice. Thus, scientists from former Czechoslovakia and the Czech Republic contributed significantly to current knowledge on *Francisella* pathogenesis and their results were highly appraised by international scientific community.

The authors would like to dedicate this review to Jiri Libich, M.D., a leading researcher on tularemia in former Czechoslovakia.

Key words: Francisella tularensis; tularemia; former Czechoslovakia; Czech Republic

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ABBREVIATIONS

 Δ , deletion;

AAA+, ATPases Associated with diverse cellular Activities:

ATP, adenosine-5'-triphosphate; dsbA, disulfide bonded protein A;

clpB, caseinolytic peptidase B; *F. tularensis*, *Francisella tularensis*; *F. tularensis* strain 15, *Francisella tularensis* subsp. *holarctica* strain 15; hsp, heat shock protein; igl, intracellular growth locus; LVS, live vaccine strain; MAPK, Mitogen-Activated Protein kinase; MHC, major histocompatibility complex; NK, natural killer; subsp., subspecies

INTRODUCTION

Background on Francisella tularensis

Francisella tularensis (F. tularensis) - a highly virulent, nonsporulating, pleomorphic, facultative intracellular, Gram-negative coccobacillus is capable of causing a zoonotic disease called tularemia in a large number of mammals. The first description of this disease was probably made on September 19, 1907 by Ancil Martin, an ophthalmologist of Phoenix, the territory of Arizona, in his letter addressed to F. G. Novy, a professor of bacteriology at the University of Michigan. Martin stated that under observation and treatment he had five cases of an infection caused by the skinning and dressing of wild jack rabbits (reviewed by [1]). The first published description of the disease may be found in the article by R. A. Pearse who reported six atypical cases of fever caused by a deer-fly bite. He called this disease a "deer-fly fever" [2]. However, it is generally accepted that the history of tularemia started in 1911 when this illness was discovered in ground squirrels in Tulare County, California by G. W. McCoy from the US Plague Laboratory in San Francisco [3]. Subsequently, the bacillus, identified as the causative agent of tularemia, was isolated and named Bacterium tularense [4]. The first human case of tularemia, which was confirmed bacteriologically, was reported by Wherry and Lamb when they isolated the bacteria from a conjunctival ulcer [5]. The man who contributed most to our knowledge of tularemia as a separate clinical entity was Edward Francis. Francis studied the "deer-fly fever" in Utah and recognized its identical traits with the illness from Tulare County and named the infection tularemia [6]. At the same time, the attention was attracted to the acute febrile disease transmitted to man by wild rabbits of the Abukuma Mountains in the eastern part of Fukushima prefecture in Japan. The illness was

known as "yato-byo" (the hare illness) or, according to the discoverer, as Ohara's illness [7]. Edward Francis and Dunlop Moore finally concluded that Ohara's disease and tularemia were identical on the basis of the exchanged clinical sample analysis in 1926 [8]. *F. tularensis* was recognized as the causative agent of "water rat-trappers' disease", an illness acquired by trappers who skinned water-rats for their pelts [9, 10]. Soon after that, tularemia was also reported in Norway (1929), Canada (1930), Sweden (1931) and Austria (1935) [11]. Several other cases of tularemia were then reported from more than 15 countries in North America, Asia, and Europe during the third decade of the twentieth century.

Taxonomy of Francisella tularensis

Tularemia is a zoonotic disease appearing in the entire north hemisphere. F. tularensis, as the etiological agent of, tularemia comprised of four subspecies: tularensis (Type A), holarctica (Type B), mediasiatica, and novicida. Genotyping methods have demonstrated that both subspecies, Type A and Type B, can be further divided into subpopulations. The subspecies differ in their pathogenicity and in their geographic distribution. F. tularensis subsp. tularensis Type A (in North America), is further divided into distinct Type A1 (east) and Type A2 (west), resp., differs according to clinical severity [12, 13], F. tularensis subsp. holarctica Type B (in North America, Europe and Asia), and F. tularensis subsp. mediasiatica (dominating in Central Asia). The taxonomy of F. novicida is still a matter of a debate. An attempt of Huber et al. [14] to classify F. novicida as the fourth subspecies of F. tularensis was refused by others [15].

Tularemia outbreaks in former Czechoslovakia and in the Czech Republic

The first outbreak of tularemia in former Czecho-slovakia was recognized in the area of south-east Moravia in 1936-1937. More than 400 cases were diagnosed during this outbreak. All affected people were in contact with hares [16]. After the Second World War, the outbreaks were mainly associated with the campaign in sugar factories. From that time, it was recognized that tularemia is endemic in the territory of Moravia as well as Bohemia. The number of reported cases is about 100 per year with some exceptions (Figure 1). The majority of cases usually appears during a hunting season. The outbreak of water-born tularemia was also registered in the area near the town of Plzen during the onset of the new millennium [17]

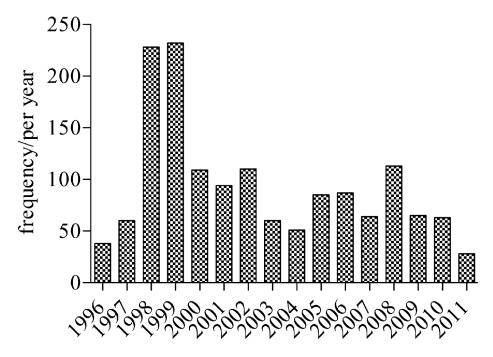


Figure 1. The amount of reported cases of tularemia in Czech Republic during the last 16 years.

The existence of tularemia outbreaks became the basis of the intensive research of this disease in the Czech Republic. Altogether three independent research groups started to study this relatively new illness in former Czechoslovakia. One group came from the Medical Faculty of Comenius University in Bratislava. This group is associated with the name of Darina Guricova. She published the first isolation of F. tularensis subsp. tularensis Type A in Europe, which was a unique report about the presence of Type A in Europe [18]. Second group was established at the branch of the Institute of Landscape Ecology of the Academy of Sciences in the town Valtice. The studies of this group were concentrated on the analyses of vectors in active enzootic foci (floodplain meadow and forest ecosystem) located mainly in South Moravia [19-21]. The third group started their research at the Military Institute of Hygiene, Epidemiology and Microbiology in Prague during the late fifties. This group was engaged solely in the experimental research, which was motivated by the political situation. The dominant person of this group was Jiri Libich, M.D., the head of the Bacteriological Division of the Military Institute of Hygiene, Epidemiology and Microbiology in Techonin. A transformed form of this research group has been working successfully till now and can be currently found at the Faculty of Military Health Sciences, University of Defence, in Hradec Kralove.

Overview of the former Czechoslovak and Czech tularemia research

Initially it was important to prepare the culture media for the cultivation of Francisellae [22] and to develop a treatment strategy for tularemia based on existing antibiotics [23, 24]. Simultaneously, an experimental model of inhalational tularemia was constructed for the study of pathogenesis and virulence of individual subspecies of F. tularensis based on several animal species. The laboratory experiments were enabled by the development of the aerosol technology and exposure procedure named Single Dose Exposure 400 (SINEX 400 for scheme see Figure 2). A great advantage of this technique was the ability to accurately calculate an inhalation and a deposition dose of microbes in lungs of experimental animals. The calculation was based on several aerosol parameters selected for the experiment (e.g. concentration of dispersed particles, stability of the aerosol, bacterial population density in particles, parameters of the aerosol chamber such as volume of the chamber, air flow in the chamber, relative humidity, temperature or the parameters of an experimental animal used for the experiments). A mathematical model of inhalational tularemia was developed based on theoretical and experimental data, and consequently was confirmed experimentally using

the SINEX 400 technique [25-27]. It was demonstrated that a number of granulomas is strictly dependent on a number of deposited microbes in lungs of infected animals, on a kinetics of microbe dissemination dependent on the

infection dose after aerosol challenge, and on generation time of both an attenuated *F. tularensis* subsp. *holarctica* strain 15 (*F. tularensis* strain 15) and a fully virulent *F. tularensis* subsp. *holarctica* strain 130 in murine peritoneal macrophages.

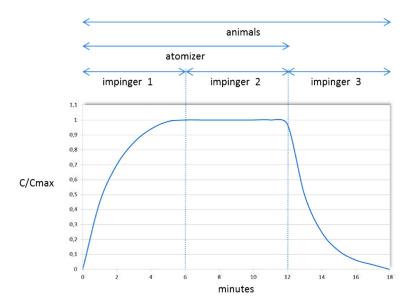


Figure 2. The scheme of an aerosol technique constructed and named by Jiri Libich as Single Dose Exposure 400 (SINEX 400). An aerosol chamber of 400 liter volume and a pneumatic nebulizer constructed for creation of monodisperse aerosol with a mean diameter of particles 3.5 um were used for an inhalation model of infection. Three impingers were put into the aerosol chamber during nebulization of 1ml bacterial suspension per minute. One impinger contains a simple cultivation medium for aerosolized bacteria. The constant flow of air was drowned through the impinger for an indicated time (axis x). A number of microbes in the aerosol was determined by plating the impinger liquid on a solid thioglycolate-glucose-blood-agar plate. An amount of bacteria in the aerosol was expressed as C/Cmax (axis y) calculated from the impinger liquid.

After this initial period, tularemia research had moved on to a study of a host immune response to experimental F. tularensis infection. The conditions for the best effectiveness of immunization against virulent strains of F. tularensis were studied during the eighties of the last century. The challenging of immunized animals with the virulent strains of subsp. holarctica Type B and tularensis Type A led to the observation that a protective effect of inhalational immunization is significantly better than a subcutaneous one. Furthermore, it was also demonstrated that the induction of protective immunity is not an exclusive property of a live vaccine. Some degree of protection was also obtained when a proliferation of a live vaccine was limited by an administration of antibiotics (streptomycin, kanamycin) in immunized animals. Similar protective effect was observed after an immunization with attenuated F. tularensis strain 15 autolysate. It was also documented that an intensity

of immune reaction and a spectrum of activated immune mechanisms were dependent on a genetic background of mice, on a route of infection, and partially on a dose of infection. A production of specific antibodies, an activation of macrophages, a blast transformation of lymphocytes, and a production of several cytokines were used for monitoring an immune response [28-30]. It was also proved that an infection with F. tularensis leads to an activation of "Natural" killer (NK) cells during early stages of infection briefly after its discovery [31]. The NK cell activation was accompanied by a production of regulatory interferon gamma cytokine [32]. Transfer experiments also confirmed an MHC class II restriction of a protective immune response induction [33]. A phenomenon of an early protective response to virulent F. tularensis Type A occurring between 24 and 48 h after immunization was published by Karen Elkins and co-workers later in 1997 [34].

Table 1. A survival of C3H/Cbi/Bom mice irradiated by 60 Co in a dose of 4 Gy. The resistance to *F. tularensis* infection is totally abrogated using gamma irradiation (60 Co) higher than 3 Gy. A significant rapid depression in a specific T lymphocyte count is observed after irradiation (3 – 4 Gy), and their nadir is reached 36 to 48 h after an irradiation event. The duration of this decrease correlates with a radiation dose, while the recovery begins 10 to 15 days after a dose of 3 – 4 Gy.

Irradiation	Infection*	Survival	%	MTD (days)
-	+	45/49	91.8	8.5
1 Gy	+	18/20	90.0	9.0
2 Gy	+	8/20	40.0	9.8
3 Gy	+	0/20	0.0	8.8
4 Gy	+	0/20	0.0	7.8
5Gy	+	0/20	0.0	6.4
5 Gy	-	20/20	100.0	-

^{*} Mice were s.c. infected with F. tularensis 15 infection in a dose of 1.05 x 10² live microbes, 72h after irradiation.

Table 2. A survival of C3H/Cbi/Bom mice irradiated by 60 Co in a dose of 4 Gy. A natural recovery of resistance to infection started during a second week after irradiation. A significant rapid depression in a specific T lymphocyte count is observed after irradiation (3 – 4 Gy), and their nadir is reached 36 to 48 h after an irradiation event. The duration of this decrease correlates with a radiation dose, while the recovery begins 10 to 15 days after a dose of 3 – 4 Gy.

Infection* after irradiation** (days)	Survival	0/0	MTD (days)	
1	0/8	0.0		
3	0/8	0.0	7.1	
7	0/8	0.0	7.0	
14	7/8	87.5	7.0	
21	8/8	100.0	-	
28	7/8	87.5	7.0	
42	8/8	100.0	-	
56	8/8	100.0	-	
Un-irradiated	15/16	93.7	8.0	
Un-infected	16/16	100.0	-	

^{*} Mice were s.c. infected with F. tularensis 15 infection in a dose of 1.05 x 102 live microbes, 72h after irradiation.

A separate set of experiments was performed using ⁶⁰Co gamma irradiation of mice as a model of immunocompromised animals. It was demonstrated that gamma irradiation of mice greater than 3 Gy totally abrogated resistance to an infection induced by *F. tularensis* strain 15 (Table 1). Minimum two weeks were required to reach full natural recovery from this deep decline of resistance (Table 2). It means that a live vaccine immunization of generally immunocompromised individuals is practically impossible. An immunization with killed microbes or microbial protein extracts lack

a sufficient protective effect. Thus, one of the possibilities how to protect irradiated individuals is a passive transfer of immunity. Both immune cells and antibodies are effective in a passive transfer of protective immunity to naive recipient (Table 3). This knowledge can conclude that specific antibodies provide some degree of protection probably mediated by an antibody dependent cell mediated bactericidal activity, originally published by Lovell at al. 1979 [35, 36]. Later, the same conclusion was published by Stephan Stenmark in 2003 [37].

^{**} Mice were infected at the indicated day after irradiation using gamma irradiation (60Co).

Table 3. A survival of C3H/Cbi/Bom mice irradiated by ⁶⁰Co in a dose of 4 Gy, passively protected by transfer of cells, sera against *F. tularensis 15* or Tularin.

Irradiation	Infection*	Treatment**	Survival	%	MTD
4 Gy	-	-	8/8	100.0	-
4 Gy	+	Tularin	0/8	0.0	8.2
4 Gy	+	Naïve spleen cells	2/22	9.1	7.9
4 Gy	+	Naïve thymus cells	1/10	10.0	7.8
4 Gy	+	Immune spleen cells	22/22	100.0	-
4 Gy	+	Immune spleen cells + NMS + C	10/10	100.0	-
4 Gy	+	Immune spleen cells + anti Thy1.2 Ab + C	10/10	100.0	-
4 Gy	+	Ultrasound destroyed immune spleen cells	2/10	20.0	8.9
4 Gy	+	Mouse serum naïve	0/10	0.0	8.6
4 Gy	+	Immune serum (3 rd day - live vaccine)	3/7	42.8	8.7
4 Gy	+	Immune serum (7 th day - live vaccine)	4/5	80.0	7.0
4 Gy	+	Immune serum (11th day - live vaccine)	10/10	100.0	-
4 Gy	+	Immune serum (21st day - live vaccine)	10/10	100.0	-
4 Gy	+	Immune serum (56th day - live vaccine)	7/7	100.0	-
4 Gy	+	Immune serum (21st day - heat inactivated vaccine)	7/7	100.0	-
-	+	-	20/22	90.9	7.5

^{*} Mice were s.c. infected with F. tularensis 15 infection in a dose of 1.05 x 10² live microbes, 72h after irradiation.

At the turn of the millennium, brand-new tularemia research stimuli were discovered on a molecular level due to the development of advanced proteomic analyses of both pathogen and infected host cells. A proteomic technology based combining various gel electrophoresis procedures, a Western blot technique, and mass spectrometry identification approaches were used for analyzing *F. tularensis* immunoreactive proteins [38-42], for identifying unique typing markers of Francisella subspecies [43-45], and for studying host-pathogen interaction at a molecular level. These studies were ranging from a subcellular proteome of bacterial membranes [46-48] to bacterial secreted proteins important for early stages of a host-pathogen interaction [49]. Highly sophisticated approaches of quantitative shotgun proteomics were then applied to a protein profiling study of bacteria exposed to stress conditions in vitro. Such conditions were able to immitate prevailing conditions inside host cells in vivo [51-54] alongside with the identification of several proteins whose expression was changed. Among

them there were proteins encoded by an igl operon, a Hsp100 chaperone ClpB with its assumed function in reactivation of aggregated proteins under in vivo stress conditions, and an ORF FTL 0200 encoding a protein of putative AAA+ ATPase of a MoxR subfamily. All these proteins seem to be indispensable for the resistance to stress conditions and are substantial factors controlling a virulence and a pathogenicity of F. tularensis. Further, the comparative shotgun proteome analyses of F. tularensis subtypes revealed several promising candidate proteins for constructing a new type of attenuated live vaccines. Deletion mutants were prepared for some of the identified proteins and two of them, FSC200 $\Delta dsbA$ and FSC200 $\Delta iglH$, were successfully tested for their attenuation and their immunogenicity. Regarding the DsbA deletion mutant, a molecular mechanism of its attenuation was studied comparing protein patterns of the original wild and deletion mutant strains. Several proteins accumulating in a membrane of the mutant strain were found and some of them were later identified as important factors of F. tularensis

^{**} Passive transfer of cells or sera was realized 2 h before infecting irradiated mice. A treatment of irradiated mice by a commercial USSR preparation Tularin (antigenic material for skin tests used frequently in the past as a prototype of dead vaccine) was realized 2 h after irradiation.

virulence [55, 56]. Current proteomic experiments are focused on structural characterization of *F. tularensis* proteins, especially on identifying bacterial membrane glycoproteins [50].

The above-mentioned proteome technologies were also used in a study of a host response to an ongoing infection. These studies clarified that a mutual interaction is stressful for both organisms. Various production of a highly stress-inducible hsp72 protein, a member of the hsp70 family, has been demonstrated in macrophages of three different inbred strains of mice exhibiting either a resistance or a susceptibility to an F. tularensis LVS infection. The hsp72 was observed to be preferentially produced and accumulated in intracellular space of a murine peritoneal adherent cell [57]. F. tularensis LVS induces apoptosis in a macrophage after infecting these cells. This process requires activation of a p42/p44 MAPK pathway and is associated with a reduced p38 MAPK activity, indicating that infection-induced cell death can be caused by perturbation of these two signalling pathways [58]. A mapping of Francisella intracellular trafficking inside macrophages discovered a complicated fate of intracellularly localized bacteria. Bacteria of F. tularensis strain LVS disintegrate a phagosome early after its entry to a host cell followed by an escape to a cytosol where it intensively proliferates. A significant part of bacterial population merges into an autophagosome expressing MHC class II molecules. Subsequently, formed autophagosome can be a source of immunogenic signal for CD4+ T cells [59].

Macrophages are considered to be primary host cells for F. tularensis, but several other cell types in the immune system also serve as host cells for tularemia infection. F. tularensis is able to infect, "disturb", and to activate not only cells of a mononuclear phagocytic system but also dendritic cells [60], epithelial cells [61] and hepatocytes [62]. Moreover, Krocova and co-workers demonstrated that F. tularensis also infects mouse (A20) or human (Ramos RA-1) B cell line cells as well as murine primary spleen B cells in series of experiments [63]. Within an infection of B cells, it has been observed that F. tularensis FSC200 activates several caspases such as caspase 8, 9 and 3. An activation of Bid, cytochrome c, apoptosisinducing mitochondria factor and proapoptic Bcl-2 family member has also been determined. This causes depolarization of a mitochondrial membrane potential in a Ramos cell line, thus leading these cells to apoptosis. Unlike live bacteria, killed F. tularensis FSC200 is capable of activating caspase 3 only and

does not cause apoptosis of Ramos cells. Killed bacteria also cause accumulation of anti-apoptotic protein Bclx_L in mitochondrial membranes. Therefore, live *F. tularensis* activates both caspase-dependent pathways (receptor-mediated and intrinsic) and caspase-independent mitochondrial death in B cells [64].

CONCLUSION

In conclusion, former Czechoslovak and consequently Czech scientists contributed significantly to current knowledge on tularemia pathogenesis and their results are highly appraised by international scientific community. Former Czechoslovak and Czech scientists underwent a long way from identifying the illness at the beginning to recent molecular analyses of a bacterial virulence during the last 50 years. Therefore, the authors would like to dedicate this article to a successful, long-time work of Jiri Libich (Figure 3), which attracted many of us to become a part of tularemia research community.

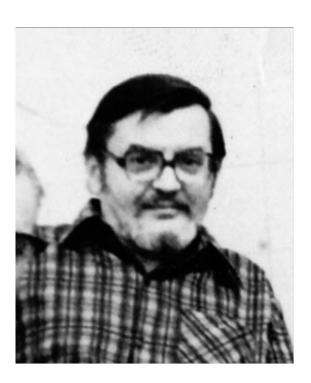


Figure 3. One of the very rare photos of Jiri Libich who was the leading researcher on tularemia in former Czechoslovakia during the sixties and the seventies during the last century.

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