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TMD Rediscovered: A New Paradigm, Part One

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LEARNING OBJECTIVES

After participating in this CE activity, the individual will learn:

- Evidence-based support for arthritis of the temporomandibular joint (TMJ) as an infectious disease.
- Elements of inflammation as a vascular event that damages blood vessels supporting the TMJ.

ABOUT THE AUTHOR



Dr. Gottesman received his dental education at New York University College of Dentistry, graduating in 1977, and currently maintains a private practice in Rockville Centre, NY. He has served on the teaching faculty of the Lenox Hill

Hospital Dental Service and has also been on the faculty of the New York University College of Dentistry TMD/orofacial pain program, where he served as director of neuropathic pain in the advanced orofacial pain/TMD graduate teaching program. Together with Dr. G. William Arnett and Dr. Stephen B. Milam, Dr. Gottesman has co-authored the acclaimed 2-part article, "Progressive Mandibular Retrusion—Idiopathic Condylar Resorption" which was published in the *American Journal of Orthodontics* in July and August of 1996. For comments, questions, or lecture information, he can be reached at lgotteswim@aol.com.

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INTRODUCTION

The human temporomandibular joint (TMJ) has been most eloquently described by Benjamin Moffett:¹ It reigns as the paladin of joints, for, having served as the evolutionary deliverer of the middle ear, it gallantly became the enduring hinge on which woman's emancipation continually swings. Its versatility fills us with respect for a structure so magnificent in function that at one moment it is the sliding pivot of trituration and mastication, at the next, the gnashing safety valve of rage and fury, and, in moments of sublime creativity, the proprioceptive junction for poetry and song.

The TMJ is a structurally unique joint positioned at the intersection of many vital cross-system interactions. Moffett's article¹ underscores the sophisticated and intricate physiologic role the TMJ plays in simultaneously coordinating multiple functions under the influence of varying functional demands.²

Much has been said about temporomandibular joint disorders (TMDs) in the dental literature. The large album of symptoms and associated diseases which seem to defy assignment to a common root cause is problematic, controversial, and confusing. The historical discourse written by Nelson and Landau³ chronicles the diverse causal heritage of TMD from 1887 forward, and features the contributions made by many professional icons of their time. No doubt, many of our esteemed colleagues have had a strong hand in shaping our professional outlook regarding TMD as a complicated, multifaceted abnormality. However, despite the many hypotheses associated with loss of normal function and degenerative joint changes, the etiopathogenesis of TMD remains unclear.

As a female gender-biased disorder, the list of symptoms and causative factors are strikingly varied, ranging from occlusal discrepancies and prematurities; occlusal pattern abnormalities; malocclusions; variations from centric relation; bruxism; jaw pain; joint pain; clicking; popping; masticatory muscle discomfort; limited range of motion; ear pain; headache; tinnitus; sore throat; neck pain; stress; anxiety; tension; psychosocial issues; tooth loss; accompaniment by back pain, autoimmune syndromes, hormonal and reproductive disorders, sleep disturbances, and gastroesophageal reflux disease, to mention a few.⁴⁻¹⁵

With such a broad band of wide-ranging associations, co-mingling, and co-morbid factors, where can practitioners look to find the foot that will fit the glass slipper?

The mystery may be solved by close examination of the key factors associated with disorders of the TMJ: cytokine profiles, inflammation, and coagulation.

Part one of this article is devoted to a supportive literary review presenting arthritis of the TMJ and other joints as an infectious disease. This portion highlights important



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elements of inflammation as a vascular event which involves activation of the coagulation system and causes damage to blood vessels supporting the joint structure and failure of hard tissues.

Part 2 will channel the academic findings discussed in the first part into the clinical setting. The second article will help prepare the clinician to identify TMD and orofacial pain patients with a comprehensive questionnaire targeting disease forms and symptoms from the position of infectious diseases and thrombophilia (pro-clotting disorders) as a mixed entity. Case studies or portions of case reports will be used to demonstrate blood abnormalities and potential blood studies which can be authorized to validate or dismiss coagulopathies and the evidentiary trail of infection. An MRI sequence prescription and clinical examination companioned with this concept will be forwarded. Part 2 will also briefly address 2 imperative questions. First, why does the gender bias exist on this platform, and second, if TMD had its origin as an occlusomuscular disorder, why would a splint work if an infection was at the core foundation?

DECODING THE MYSTERY

Cytokines

Cytokines are proteins that serve as signaling molecules, some of which have pro-inflammatory properties, such as tumor necrosis factor-alpha (TNF- α), interleukin-1 beta (IL-1 β), IL-6, interferon gamma, granulocyte-macrophage colony-stimulating factor (GM-CSF), and IL-8, to name the most commonly cited. Pro-inflammatory cytokines have been implicated in joint pain, TMJ closed lock, inflammation, internal disc derangements,¹⁶⁻²² and joint effusions.²³⁻²⁷ Other cytokines are anti-inflammatory, including IL-1 receptor antagonist, IL-4, IL-6, IL-10, IL-11, and IL-13,²⁸ and may reflect an immunomodulatory dampening counterbalance to the cytotoxic response precipitated by the pro-inflammatory group.²⁹

Cytokine profiles in asymptomatic controls differ from abnormal TMJ patients.³⁰ Both pro- and anti-inflammatory cytokines are frequently found together in symptomatic patients.^{22,31,32} Additionally, several members of this signaling family have dual roles in modulating the inflammatory immune response and other regulatory processes as demonstrated by IL-6.^{32,33-37} Cytokines maintain a fundamental role as part of the innate immune system, serving as acute phase reactants in the initial nonspecific response to infection. The innate immune cells comprise a population of predominantly white blood cells (WBCs) forming the frontline of defense when our barrier system(s) have been breached. Cytokines are released by resident members of this WBC lineage, including neutrophils, natural killer cells, dendritic cells, monocytes, basophils, eosinophils, and local tissue mast cells and macrophages.³⁸⁻⁴⁴ As part of a complicated and orchestrated system, the cytokines are responsible for immune cell recruitment and other cell-mediated trafficking functions.⁴⁵⁻⁴⁷

Pro-inflammatory cytokines, free radicals, and oxidized low-density lipoprotein are among the molecules that can activate nuclear factor kappa-light-chain-enhancer of activated B cells (NF-k β). The receptor activator of NF-k β , called RANK, is active in bone regulation, and upon exposure to cell-mediated pro-inflammatory cytokines can initiate osteoclastogenesis.⁴⁸

The key element, however, in understanding the character of cytokines is that *they are most commonly activated in response to microbial pathogens.*⁴⁹ Therefore, part of the differential diagnosis in the pathophysiology of TMD must include infection, a lexicon missing from our current common wisdom.

Central to the theme of inflammation is activation of the coagulation system. Triggered by the cytokine release in response to bacterial, viral, fungal, and parasitic organisms, activation of the clotting cascade ensues with the production of tissue factor (thrombin formation) and induces a local or more global systemic vascular response, resulting in a hypercoagulable state,49 whereby the blood becomes thickened (more viscous). In 1856 the German physician, Rudolf Virchow, described the trio of damage to the blood vessels and changes in blood viscosity leading to this hemostatic abnormality called Virchow's Triad: vascular endothelial injury, blood stasis, and hypercoagulability of the blood. This phenomenon contributes to the domino effect of disturbed flood flow, impaired oxygen delivery to tissues, and hypoxia, and raises the risk for venous thromboembolism or deep vein thrombosis.50,51 At the tissue level, larger vessels that become obstructed by



thrombus formation may give rise to emboli, which can have catastrophic consequences such as a pulmonary embolism, myocardial infarction, or stroke. However, medium and small portions of the vascular system can also be adversely affected, leading to local ischemic changes and peripheral nerve involvement as the bloodbreached. nerve barrier is compromising the blood supply to sensory and motor nerves with resultant motor and painful sensory neuropathies.52,53



Figure 1. The relationship among microbial pathogens, cytokines, inflammation, and coagulation.

Further, the activation of the hemostatic system concomitant with inflammatory changes leads to fibrin generation with the formation of fibrous tissue in both intravascular and extravascular fluid spaces.54 Patients with synovitis,55-57 rheumatoid arthritis,58,59 adhesions,60 and the consequential development of free radicals⁶¹ demonstrate fibrin production in association with joint abnormalities and other inflammatory diseases. As the endothelium of the blood vessel, particularly the venous component, becomes paved and thickened with fibrin, its lumen caliper narrows, partially or fully obstructing the venous outflow. Changes in the integrity of the vascular wall lead to arteriosclerosis (stiffening of the blood vessel wall) and decreased compliance.62 These sequelae set the stage, creating the potential for a static backflow system with an increase in venous pressure and reduced arteriole blood flow influx. With changes in blood flow rate, vascular endothelial wall shear stresses are altered. Adverse modifications in hemodynamics escort and enhance fibrin generation on the venous end, while the arteriolar transformation favors a platelet-rich environment which will ultimately be enriched by the incorporation of fibrin strands into the platelet aggregate.54,62

Such abnormalities in blood flow can have profound and dramatic consequences for a joint that is literally being suffocated to death; the accompanying disturbances are implicated in the development of bone marrow lesions⁶³ and osteonecrosis.⁶⁴

In 2011, Jennewein et al⁶⁵ stated that: "Coagulation is fundamental for the confinement of infection and/or the inflammatory response to a limited area. Under pathological inflammatory conditions such as arthritis, multiple sclerosis, or sepsis, an uncontrolled activation of the coagulation system contributes to inflammation, microvascular failure, and organ dysfunction."

The purpose of coagulation is to limit infection and enhance the inflammatory response with the hope of immune functions isolating and quelling an infectious or injurious event.^{66,67} The intimacy of cross-talk between systems is essential in trying to strike a balance that addresses the diseased state. The relationship between the severity of the inflammatory reaction and the rigor of the hemostatic response may be beneficial, but not without consequence.⁶⁸ Despite what we might insinuate is nature's purpose, the cytokines and other acute phase reactants which are first responders in an attempt to kill microbial trespassers⁴⁴ have a highly cytotoxic effect on both the micro-organisms and resident tissues. Ultimately, human body parts caught in the crossfire summarily fall victim to "friendly" fire (Figure 1).

While much attention has been devoted to the biomechanical aspects of joint loads and the influence forces have on the local structures as a rationale for TMD,⁶⁹⁻⁷¹ the cytokine profiles in joint studies which have explored the potential for pro-inflammatory cytokine production during cyclic compressive forces are disparate



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as demonstrated by Muroi et al in 2007,72 whereby TMJ synovial tissue surgical explants exposed to mechanical compressive forces produced mostly matrix metalloproteinases along with the chemokine IL-8, without the expression of TNF- α , IL-1 β , IL-6 in their recordable assays. At this juncture, the lack of TNF- α , IL-1 β , and IL-6 cytokine expression during mechanical compressive loading becomes a very important issue. If these cytokines have a primary allegiance to the innate immune system, but are not produced during load testing studies, then it raises the likelihood that part of the root cause for TMD arises from infections. The failure of the joint as an organ could occur at multiple tissue levels73 without the introduction of mechanical overload and could be potentiated by persistent clotting abnormalities introduced by temporary or persistent microbial invaders.49

Orhan et al⁷⁴ have demonstrated, via MRI, TMJ bone marrow shifts in patients with anemia without the requirement for disc displacement or internal derangement. The prospect that inherited intrinsic genetic risk factors such as thrombophilia (pro-clotting disorders) may precipitously amplify the clotting cascade during infection must also be considered.⁶⁴

Is TMD an Infectious Disease?

The literature supporting TMD as an infectious disease is largely succedaneous and reactive to the greater number of articles written by researchers within the medical profession. However, the lesser body of evidence in the dental space demonstrating the presence of pathogenic organisms either in the peripheral venous blood, or TMJ tissues or fluid, warrant merit and lend credence to the concept that infection is involved with TMJ disturbances, and may be the leading cause. Although many of the dental articles are highly conserved in the pursuit and identification of only singular microbial species, there are a few studies which utilize a more broadband analysis in parallel with the medical writings to demonstrate co-infection or polymicrobial diversity related to TMJ disease.

Chlamydia trachomatis, one of the organisms responsible for sexually transmitted diseases, occupies a very large presence in both the medical and dental literature and is classified symptomatically under the umbrella of reactive arthritis. Henry et al^{75,76} have demonstrated the presence of C trachomatis in internal derangement surgical studies where the peripheral blood analysis was positive for immunoglobulin antibodies demonstrating cases of both past and active infections. In a more comprehensive search, Henry et al^{75,76} expanded their investigation beyond C trachomatis, and included Mycoplasma fermentans and orale, Mycoplasma genitalium, and the organisms associated with food poisoning, Campylobacter iejuni spp., Yersinia enterocolitica spp., Salmonella spp., and Shigella spp.⁷⁷ The latter group has also been included under the auspices of reactive arthritis. Surgical tissue samples were harvested from the bilaminar zone of 26 surgical patients. Of the 26 patients (24 female and 2 male), 42% had C trachomatis, 35% M genitalium, and 23% M fermentans/orale. The food poisoning group was negative in this study. Eight percent of the group tested positive for 2 organisms (M fermentans/orale and M genitalium), while 4% were positive for a combined cluster of 3 organisms (C trachomatis, M fermentans/orale, and M genitalium).78 In 1999 and 2007, Henry et al^{76,79} conducted more constrained studies, looking only for C trachomatis and serovars from the bilaminar zone of surgical specimens. Of 70 patients in the 2007 study, 46% were positive for *C trachomatis*.⁷⁶ Further, in the same investigation, Henry et al⁷⁶ found differences in the cytokine profiles between symptomatic Chlamydia positive patients and the control group, showing elevated levels of TNF- α and IL-6 in the symptomatic population tested for these cytokines.

Kim et al⁷⁷ in 2003 performed a relatively expanded synovial fluid study and found the presence of *M genitalium* in 86% of the 43 patient samples with a 1:1.69 male to female ratio. Of this group, 33 symptomatic participants had a displaced disc, and 10 participants were asymptomatic nondisc displacement recruits. The other organisms investigated were *Staphylococcus Aureus*, present in 52% of the population tested; *M fermentans/orale* 37.2%; *Actinobacillus actinomycetemcomitans* (a highly pathogenic periodontal disease-related species) 25.6%; and *Streptococcus mitis* 7.0%. The conclusions drawn predicated on the higher levels of *S Aureus* present in the symptomatic displaced disc group versus the non-displaced disc asymptomatic volunteers was





that S Aureus was implicated as causal to TMD. 77

Adachi⁸⁰ shows the involvement of *Mycobacterium* in the TMJ. Sun et al⁸¹ provide a broadband peak cell rate (PCR), 16s ribosomal RNA analysis of synovial fluid confirming the presence of 11 species and substantiating that a wide variety of bacteria can be associated with TMD. Other authors have made significant contributions to this area as well.⁸²⁻⁸⁴ Jeon et al⁸⁵ have identified that the source of infection can be hematogenous with respect to the TMJ, and corroborates a statistical correlation among TMJ disease, sinusitis, rhinitis, pharyngitis, and tonsillitis.⁸⁶

In order to help consolidate and provide a clinical perspective, the addition of some of the comparable medical literature may prove

helpful. In recapitulating what has been stated thus far, some salient points which need to be placed into context will be reviewed.

First, there is a regional flavor for the path of origination and dissemination to the TMJ from neighboring structures. Sinuses, nasal passages, pharynx, upper respiratory, and tonsils are all in close proximity to the TMJ. Infections in these nearby areas are potential origination sites for resident organisms to gain migratory access through the vascular or lymphatic systems in order to reach and occupy a niche environment.^{85,86}

Secondly, the oral cavity is a substantial reservoir for micro-organisms. The study by Kim et al⁷⁷ demonstrates involvement of the periodontal pathogenic organism *A actinomycetemcomitans*. However, to strengthen the case for the involvement and presence of oral pathogens in joint diseases, the work of Moen et al^{87,88} validates the oral-systemic link in identifying the DNA from oral bacteria in the synovial fluid and serum from patients with rheumatoid, psoriatic, and osteoarthritis. Moen et al^{87,88} found greater diversity and concentrations of DNA from oral micro-organisms in the synovial fluid than the serum. Both rheumatoid and psoriatic arthritis had greater bacterial variety and content when compared to osteoarthritis. There



Figure 2. Symptoms of reactive arthritis.

was some disease-related specificity with respect to the organisms present and associated arthritic form. Moen et al^{87,88} suggest that the synovial tissues are a preferred niche environment for many organisms; perhaps due to their rich vascularity, presence of a basement membrane, and anatomical form, joints are an attractive sink-hole for colonization.⁸⁹

Blood serum has reduced diversity and numbers of bacteria compared to synovial tissues and fluid. This can be problematic from an investigative standpoint when performing blood studies because the immune response mounted in the peripheral blood often becomes pale in comparison to inflammatory changes in the joint. The patient may not manifest with blood draw biomarker abnormalities due to this phenomenon. The confirmation of this difficulty is established by Lyme polymerase chain reaction studies with analysis from different sites. Comparing blood, cerebrospinal fluid, synovial fluid, and skin tissue, positive markers were significantly higher in synovial fluid and tissue.⁹⁰ Also, it is important to mention that most micro-organisms have evolved evasive strategies to avoid immune detection, making positive identification difficult.⁹¹⁻⁹³

Further, organisms can migrate from sites distant from the initiating source of the infection, to joints and other sites.



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This is particularly true of the organisms associated with sexually transmitted diseases⁸⁹ and the *Mycoplasma* group.⁹⁴ Organisms included in the reactive arthritis group may originate from the urogenital tract, or gut, as in the food poisoning group, associated with *Campylobacter jejuni* spp., *Yersinia enterocolitica* spp., *Salmonella* spp., and *Shigella* spp.⁹⁵

Microbial pathogens may also entertain occupancy in tissues other than joints, finding their way to mucosa-associated lymphoid tissue as is the case with Helicobacter pylori in oral glandular tissue and its contribution to Sjögren's disease symptoms.96 Most importantly, while on their pilgrimage, organisms may affect many tissues, causing considerable and sometimes serious disturbances. What we may have considered as referred pain patterns and loosely associated disorders more accurately reflects the effect of pathogenic tenancy with genetic and metabolic alterations at multiple locations. The expressions of seemingly unrelated symptoms often seen in association with TMD are more likely to have a common denominator, under the wings of this platform, than previously thought.

This concept is manifest in the reactive arthritis symptom profiles (Figure 2 [symptoms of reactive arthritis]; Table [organisms associated with "infectious" reactive arthritis]).

Cox et al⁹⁷ and Kempsell et al⁹⁸ provide valuable information in investigating the viability of bacterial organisms in synovial tissue samples from arthritic patients. Using a reverse transcriptase-PCR analysis of bacterial ribosomal RNA, they were able to detect and characterize the bacterial diversity and existence of live organisms associated with the arthritides.^{97,98} They make a significant contribution to our understanding of joint diseases. In the Kempsell et al study,⁹⁸ 23 participants were evaluated. A comparative bacterial analysis was performed between rheumatoid arthritis (RA) and non-RA

patients. Nine patients were classified as having RA. In the non-RA group, the breakdown was as follows: 7 patients had osteoarthritis (OA), 4 patients had undifferentiated

Table. Organism	s Commonly Cited	as Associated With	Infectious or Rea	active Arthritis	
GENITOURINARY	GITRACT	RESPIRATORY TRACT	VECTOR-BORNE	ORAL/MIXED	
STD	SHIGELLA Flexneri	MYCOPLASMA	LYME DISEASE	MYCOPLASMA	
CHLAMYDIA Trachomatis	YERSINIA Enterocolitica, Pseudotuberculosis, Sonnei	Pneumoniae	RICKETTSIA	• Fermentans • Salivarium	
NEISSERIA Gonorrhoeae	SALMONELLA Entiritidis, Typhimurim	CHLAMYDIA Pneumoniae, Psittaci	ANAPLASMA	P gingivalis Treponema Denticola A actinomy-	
TREPONEMA Pallidum	CAMPYLOBACTER Jejuni Shigella, Yersinia, Salmonella, and Campylobacter = food poisoning group.	STAPH, Aureus	BABESIA	cetemcomitans PREVOTELLA (Nigrescens, Intermedia) B forsynthia	
				BIOFILM	
RHEUMATOID Arthritis (RA) Reactive Arthritis	H PYLORI	PSEUDOMONAS Aeurginosa	BARTONELLA	FUNGI	
MYCOPLASMA	E COLI	STREP, Pyogenes, B-hemolytic	EHRLICHIA	PROTOZOA	
• Fermentans	C DIFFICILE		COLORADO TICK FEVER	UNKNOWN Organisms	
• Genitalium	TOXOPLASMA GONDII	SKIN	Q FEVER		
• Homonis	ENTAMOEBA HISTOLYTICA	STAPH, <i>Epidermis</i> <i>P</i> Acnes MRSA	ROCKY MOUNTAIN SPOTTED FEVER		
	GIARDIA LAMBLIA				
UREAPLASMA	BRUCELLA ABORTUS	VIRAL			
• Urealyticum	T Whipplei	HIV			
		MEASLES			
		HHV			
		EBV			
		CYTOMEGALOVIRUS			
		B19			

Note: The term *infectious arthritis* within the context of this article is used to describe the association of microbial pathogens with autoimmune disorders like RA, as an infectious process affecting joints and other tissue structures.

Combined adaptation from references 1 to 3.

 Martinez-Martinez RE, Abud-Mendoza C, Patiño-Marin N, Rizo-Rodríguez JC, Little JW, Loyola-Rodríguez JP. Detection of periodontal bacterial DNA in serum and synovial fluid in refractory rheumatoid arthritis patients. *Journal of Clinical Periodontology*, 2009;36(12):1004-1010.

2. Ogrendik M. Rheumatoid arthritis is linked to oral bacteria: etiological association. *Mod Rheumatol.* 2009;19(5):453-456.

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arthritis (UA), and 3 patients were normal (no arthritis). Collectively, synovial tissue samples from the RA group showed the presence of 92 living individual species



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compared to 50 species in the non-RA control group. Overall, the 2 groups shared 21 species. The normal group had no organisms (Figure 3)!

The following is a synopsis of the article outcome:

• The RA group had a greater number of species and bacterial load than the non-RA group.

• There were many species unique to the individual groups and many organisms.

• There were many unidentified novel organisms which did not conform to any data sets of known organisms.

• Previously thought to be sterile, the joints in diseased patients contain multiple bacterial species.

• These organisms are difficult to cultivate or are culture-resistant, owing to the requirement of unknown fastidious culture mediums or sensitivity to osmotic gradients consistent with cell-wall deficient (CWD) bacteria.

• The colonization suggested these organisms could live in a biofilm.

• There was no uniformity with respect to synovial tissue colonization.

• Not all parts of the same tissue carry the same bacterial load or organisms.

• Microcolonization of different tissue areas is likely.

•Organisms could occupy both intracellular and extracellular locations.

• Commensals from the gut and skin were frequently found in joints and suggested trafficking from these sights.

• The presence of biofilm may confer antibiotic resistance and therapeutic challenges.

• These infections are often slow growing and can take many years to mature.

The findings by these 2 authors and others confirm the realities confronting our profession in the diagnosis and etiology of TMD. Cox et al⁹⁷ and Kempsell et al⁹⁸ have authenticated that even in the case of classical osteoarthritis, which was thought to be related to the biomechanical "wear and tear" of a joint, significant amounts of bacterial organisms are present. Olmez et al⁹⁹



Figure 3. Rheumatoid arthritis (RA) and non-RA subjects shared 21 species of microbial organisms.

recapitulates this theme in OA patients with *C trachomatis* and *Chlamydia pneumoniae*.

CONCLUSION

Pivotal to the supportive arguments presented are the thought processes that should accompany the clinician's diagnostic investigations. Inflammation and coagulation are interdependently linked, especially when infection is the precipitating source. This discourse brings together several elements which have not been brought to the general attention of the dental profession as causal to TMD. Pathogenic microbes that migrate to, and homestead, a particular site will violate the security of our defense systems, activate the release of inflammatory cytokines, and trigger the clotting cascade. These series of events, as part of the inflammatory response, will cause endothelial dysfunction, disturbed blood flow, and oxygen impairment, and lead to the breakdown of the TMJ on multiple levels.

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POST EXAMINATION QUESTIONS

- 1. Cytokines that have pro-inflammatory properties include:
 - a. Tumor necrosis factor-alpha (TNF- α).
 - b. Interleukin-1 beta (IL-18).
 - c. IL-4.
 - d. Both a and b.

- 2. Cytokines are proteins that serve as signaling molecules. Both pro- and anti-inflammatory cytokines are frequently found together in symptomatic TMJ patients.
 - a. The first statement is true, the second is false.
 - b. The first statement is false, the second is true.
 - c. Both statements are true.
 - d. Both statements are false.
- 3. Cytokines are most commonly activated in response to microbial pathogens.
 - a. True.
 - b. False.
- 4. Cytokines maintain a fundamental role as part of the innate immune system. Cytokines serve as acute phase reactants in the initial nonspecific response to infection.
 - a. The first statement is true, the second is false.
 - b. The first statement is false, the second is true.
 - c. Both statements are true.
 - d. Both statements are false.
- 5. Under pathological inflammatory conditions such as arthritis, multiple sclerosis, or sepsis, an uncontrolled activation of ______ contributes to inflammation, microvascular failure, and organ dysfunction.
 - a. IL-4.
 - b. The coagulation system.
 - c. IL-6.
 - d. All of the above.
- 6. Research by Muroi found that mechanical compressive loading of the TMJ produced mostly:
 - a. IL-1 β cytokine.
 - b. IL-6 cytokine.
 - c. Matrix metalloproteinases and IL-8 cytokine.
 - d. TNF- α cytokine.
- 7. Research has found a statistical correlation between TMJ disease and:
 - a. Sinusitis.
 - b. Rhinitis.
 - c. Pharyngitis and tonsillitis.
 - d. All of the above.



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- 8. Blood serum has reduced diversity and numbers of bacteria compared to synovial fluid.
 - a. True.
 - b. False.
- 9. Microbial organisms can migrate from sites distant from the initiating source of infection, to joints and other sites. This is particularly true of organisms associated with sexually transmitted diseases.
 - a. The first statement is true, the second is false.
 - b. The first statement is false, the second is true.
 - c. Both statements are true.
 - d. Both statements are false.

- 10. Kempsell et al's study of patients with rheumatoid arthritis (RA) found that:
 - a. The RA group had fewer number of microbial species than the non-RA group.
 - b. There was uniformity with respect to synovial tissue colonization by microbial species.
 - c. Previously thought to be sterile, joints in diseased patients contain multiple bacterial species.
 - d. None of the above.



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