NMT UC Enhanced Viral Cascade Pathway (EVCP)

H-MB = human mind-body

M-MB = microbial mind-body

H/M-MB = human/microbial mind-body = H/M mind-body

1) TFS (transformed-from state) and TTS (transformed-to states) of the system:

a) **The TFS (transformed-from system state):** Create a brief statement that describes the subject system in terms of current undesirable conditions and behaviors. Emphasize system characteristics related to the subject matter of this NMT clinical pathway. The subject system is usually the mind-body of the person we are working with, but the concepts of any NMT pathway may be applied as a lens through which to observe and transform any system, e.g., we might view the conditions and behavior of some corporation or business through the lens of any NMT clinical pathway. Static MRT (muscle response test) for the total system burden (TSB) that this state seems to the body as a $\log_{10}1000$ metric.

b) **The TTS (transformed-to system state):** Create a brief statement that describes the subject system in an idealized, perfected, healed, and transformed state, with emphasis on characteristics related to the subject matter of this NMT clinical pathway. Static MRT for the TSB (total system burden) that this state seems to the body as a $\log_{10}1000$ metric.

c) Apply the following UCS (universal correction statement) with the intention to resolve any IFs (information faults) that may constitute an obstacle to manifesting the intended transformation: **"Shall the H/M-MB now: Project its awareness, attention, belief, and intention into all times and dimensions of the undivided universal field of consciousness and identify, accept, and release attunement to all**

events, IFs, and IFs-h (IFs that hide subject IFs from awareness) that compromise awareness of, or define into existence, the transformed-from system state? <MRT-S-a>, "Or, that cause this transformation to seem dangerous, undeserved, or impossible to achieve? <MRT-S-a>, "Create awareness, acceptance, attunement, and belief of all corrective events and TrIQs (transformative information quanta) to define into existence the transformed-to system state? <MRT-S-a>, "Create the intention to collapse the quantum potential of the field's reality quanta to manifest this transformation? <MRT-S-a>, "Create optimal attunement and coherence to the source of creation, innate intelligence, and the consciousness of healing and accept the role of co-creator of all experienced reality, NOW?" <MRT-D-c>,

2) **"Shall the H/M-MB now: Project its awareness, attention, belief, and intention into all times and dimensions of the undivided universal field of consciousness and efficiently survey subject system conditions related to the following queries? <MRT-S-a>, "Identify, accept, and release attunement to all events, IFs, and IFs-h** (IFs that hide subject IFs from awareness) **that define into existence the transformed-from system state? <MRT-S-a>, "Or, that cause this transformation to seem dangerous, undeserved, or impossible to achieve? <MRT-S-a>, "Create awareness, acceptance, attunement, and belief of all corrective events and TrIQs to define into existence the transformed-to system state? <MRT-S-a>, "Create the intention to collapse the quantum potential of the field's reality quanta to manifest this transformation? <MRT-S-a>, "Create optimal attunement and coherence to the source of creation, innate intelligence, and the consciousness of healing and accept the role of co-creator of all experienced reality, NOW?" <MRT-D-c>,** Hold this MRT intention for each section that follows. Refresh this awareness and intention with each MRT pulse. The dynamic MRT end point signals that the survey is complete, awareness is optimal, and this transformation is to be informationally realized.

“Shall the H/M-MB now allow optimal awareness and correction of the following conditions with attention to CNS, Autonomous NS, and cell and cell network information processing?”:

(autonomous N.S. = Sympathetic N.S., Parasympathetic N.S., Enteric N.S., Autonomous N.S. of the spinal cord, Autonomous N.S. of the heart, Lymphatic-Immune Information Processing System)

a) **“Compromised H/M-MB attunement and awareness of predatory pathogens in the local or distant environment with attention to hemorrhagic viruses, flus, mycoplasma, spirochetes, stealth pathogens, coronaviruses, and other dangerous microbes?” <MRT-S-a>.** (Reference organisms of interest, regions and hosts where the pathogen of interest originates, vectors of movement of the pathogen - airplane travel, new locations or populations experienced with the pathogen infection of interest, e.g., hospitals, schools, responders, etc. MRT for and use the EVCP Worksheet to record desired metrics.)

b) **“Any compromise of optimal host cellular awareness of the presence in cell cytoplasm of receptor/signaling proteins positioned to recognize, link to, and tag pathogen chemistries for transport to and from the cell nucleus for analysis and to coordinate production of immune modulators and signaling chemistries for an optimal immune response?” <MRT-S-a>, Cell potential to initiate and regulate production of hundreds of interferons, cytokines, and other immune response proteins to eliminate infection, regulate inflammation, and to trigger apoptosis of viral and other pathogen infected cells that endanger the cellular community?” MRT-S-a,**

c) **“Bradykinin, an endothelium-dependent vasodilator and motor signaling agent to non-vascular smooth muscle in the bronchus and gut, increases vascular permeability, and lowers the sensitivity threshold for pain perception contributing to allodynia?” <MRT-S-a>, “Bradykinin role in inflammation, and released locally from mast cells and basophils during tissue damage?” <MRT-S-a>, “The bradykinin receptor B1, expressed from endothelial cells, only as a result of tissue injury, and is presumed to play a role**

in chronic pain, and constitutionally distributed B2 vasodilatory receptor?" <MRT-S-a>, "The role of bradykinin as a provocatory of the cytokine storm response to infectious tissue injury including that from viruses or other pathogens?" <MRT-S-a>, "Optimize the physiology of bradykinin to efficient hemodynamics, pain perception, and well-modulated humoral immune response to infection, physical injury, or other provocateurs of inflammation?" <MRT-D-c>,

d) **"Any compromise of full, balanced, and timely awareness of the arms of the H/M-MB immune response to cellular and viral pathogens?" <MRT-S-a>, Innate cellular immunity, immediate, less specific, triggered by nucleus and mitochondrial awareness of cell membrane breach by toxin, virus, or other pathogen, and preceding the onset of acquired immunity by 4-7 days?" <MRT-S-a>, "Potential recruitment of innate immunity by free ATP and/or immune cell signaling of the innate immune response to the tissue, organ, system, and whole-body level, in which successful innate response produces negative feedback to the acquired immune system preventing the unnecessarily aggressive cytokine storm response?" <MRT-S-a>, "Acquired immunity, consisting of the myeloid arm of phagocytic granulocytes, macrophages, assisted in targeting by natural killer cells, helper cells, and dendritic cells, and the histaminic mast cells controlling fluid dynamics in the tissues?" <MRT-S-a>, The lymphoid arm consisting of B lymphocytes, differentiating on activation into plasma cells that secrete antibodies; and cytotoxic T lymphocytes, which kill cells infected with viruses, and a class of T cells that activate other cells such as B cells and macrophages?" <MRT-S-a>, "Optimize an integrated and balanced response of all components of immune response, exploiting the potential of a successful initial innate response to diminish and even eliminate the need for an acquired immune response to any toxin or pathogen?" <MRT-D-c>,**

e) **"The potential of flus, hemorrhagic viruses, and other infections to block the entry and accumulation of receptor/signaling proteins in the cell nucleus needed to calculate cell immune response, to make counterfeit signaling proteins and other chemistries that**

prevent or delay cell immune response and confuse intercellular communication producing a poorly coordinated response to infection, and to hijack cell chemical production resources to replicate microbial DNA, RNA, proteins, enzymes, and viruses?" MRT-S-a, "Any such conditions that prevent, delay, or confuse efficient cell signaling and coordinated immune response within cells, tissues, and the neuroimmune system?" MRT-S-a, "Any current microbial hijacking of cell resources for chemical synthesis that furthers pathogen reproduction and spread of infection?" MRT-S-a, "Optimize cellular awareness and response to recognize and eliminate pathogen cooptation of cell resources and promote efficient cell metabolism, immune response, and cell signaling of this danger to the whole cellular community?" <MRT-D-c>, (This mechanism occurs with Ebola, Marburg, flus, and other pathogens.)

f) **"Cellular capacity to recognize the morphological, chemical, and energetic distinctions between viral and microbial counterfeits and legitimate cell signaling and immune modulating chemistries?" <MRT-S-a>, "Cellular capacity to block binding of counterfeit immune modulators to cell cytoplasm receptor proteins by energetic means and chemically by producing inert receptor agonists to host cell viral binding sites?" <MRT-S-a>, "Cellular capacity to enhance cell cytoplasm receptor protein sensitivity and selectivity to pathogen chemistries and to enhance the cell nucleus immune response, with attention to the adaptive cell danger response oxidative metabolic cycle, and mechanisms to quarantine incursions of viruses and toxins within the cell for destruction and to block new incursions?" <MRT-S-a>, "Optimize cell intelligence and awareness to distinguish microbial counterfeits from legitimate cellular immune modulators and signaling proteins, to chemically and energetically silence and/or resist such chemical confusion to produce a timely, measured, and efficient cellular innate immune and H/M-MB neuroimmune response to viral or other pathogens?" <MRT-D-c>, "Optimize adaptive cell danger response (CDR) to produce superoxides, hydrogen peroxide, and lytic enzymes to destroy any invading viral infection and related toxicity, to attract an efficient neuroimmune response, to lock down the cell**

membrane restricting chemical movement into and out of the cell, to create conditions of oxidative shielding of host cell protein building apparatus to deprive viral and other pathogen cooptation of these resources to replicate the virus or produce other proteins that aid the pathogen, and to immediately terminate the CDR cycle and return to normal cell metabolic conditions on completion of these protective operations?" <MRT-D-c>,

g) **"Awareness of the renin–angiotensin–aldosterone system (RAAS) of hormones that regulates blood pressure, fluid and electrolyte balance, vasoconstriction, and vasodilation?" <MRT-S-a>, "Production of aldosterone, the main mineralocorticoid steroid hormone from zona glomerulosa of the adrenal cortex for sodium conservation in the kidney, salivary glands, sweat glands, colon, and lungs?" <MRT-S-a>, "The activation of RAAS when carotid baroreceptors or kidney juxtaglomerular cells detect hypotension or reduced renal blood flow triggering the enzymatic conversion of renin in the blood from storage form to active form?" <MRT-S-a>, "The action of active form renin on a liver produced precursor protein to create inactive angiotensin I and its conversion to vasoconstrictive angiotensin II by angiotensin-converting enzyme (ACE) ?" <MRT-S-a>, "The presence of ACE as a transmembrane protein on vascular endothelium and organ cells with attention to capillary beds of the lungs, heart, kidneys?" <MRT-S-a>, "The presence of angiotensin converting enzyme 2 (ACE2) on outer cell membranes of cells in lungs, arteries, heart, kidney, and GI-biliary tissues that converts vasoconstrictor angiotensin II into vasodilator angiotensin 1-7 in opposition to the effect of ACE?" <MRT-S-a>, "The role of angiotensin 1-7 to improve lung hemodynamics and oxygen saturation and to reduce lung damage and risk of pneumonia in virus induced severe acute respiratory syndrome (SARS) and acute respiratory distress syndrome (ARDS) and in bacterial lipopolysaccharide-induced ARDS, and to produce similar protection to other organs and vessels?" <MRT-S-a>, "Awareness of failed RAAS control in any vessels, tissues, or organs with attention to mucous and serous membrane structures resulting in vascular hemorrhage, vascular**

effusion, and interstitial or alveolar edema of the lungs and other organs?" <MRT-S-a>, "Optimize management of the RAAS with attention to conditions of infection and inflammation to most efficiently control local and systemic blood pressure, fluid and electrolyte balance, vasoconstriction, and vasodilation to preserve and restore lung, heart, kidney, gut, vascular or other tissues under attack?" <MRT-D-c>,

h) "Any compromise of H/M-MB awareness regarding cellular production and distribution of cytokines and interferons to mobilize immune cells into areas of injury, infection, or metaplasia in a timely, efficient, and proportional manner?" <MRT-S-a>, "Compromised modulation of tissue inflammation and congestion to decrease tissue spread of pathogens, and to initiate appropriate fever to enhance the rate of metabolic immune activity?" <MRT-S-a>, "Any confusion, misperception, or incomplete awareness of neuroimmune or cellular intelligence that may contribute to positive feedback conditions that fuel a cytokine storm of violent, inappropriate, and untimely cytokine production and resultant inflammation, tissue-injurious inflammation intensity and fever, failed control of the volume and chemical makeup of body fluids and gels with attention to decreased fluid and gel viscosity favoring pathogen mobility – with attention to intracellular, extracellular, lymph, blood, CSF, mucous, serous, and interstitial fluids and gels?" <MRT-S-a>, "Virus induced cytokine triggering of mitochondrial death factors that kill bystander lymphocytes and compromise effective immune response?" <MRT-S-a>, "Manifest cell awareness and capacity to recognize, destroy, or silence counterfeit viral immune modulators and signaling chemistries and to optimize cell awareness and chemical and energetic signaling to create timely, proportional, and efficient cellular metabolism, fluid systems volume and chemical makeup, and efficient innate cellular and neuroimmune response?" <MRT-D-c>, "Manifest cell awareness and capacity to recognize structural chemical similarities of any pathogen of current interest to historical pathogens to which the H/M mind-body has mounted successful defenses and produce antibodies

to attack such revealed vulnerabilities and to modify cell receptors structures and populations to minimize the infectious potential of pathogens?" <MRT-D-c>

i) "Any compromise of H/M-MB awareness of virus-cell attachment mechanisms: Hemorrhagic virus, flu virus, or other pathogen glycoproteins introduced to cell membranes to compromise membrane strength and strong cell junction adhesions required to resist spread of the pathogens through the tissues?" <MRT-S-a>, "The presence of many decoy epitope binding sites on hemorrhagic virus glycoproteins that snare the cell's virus neutralizing cellular antibodies, attach them to cell membrane and so promote virus-cell attachment-infection?" <MRT-S-a>, "Hemorrhagic virus, flu virus production of hemagglutinin "glue" molecules that bond viruses to epithelial cells of serous and mucous membranes and endothelial vessel cells to infect respiratory, urinary, neurological, endocrine, vascular, or lymphatic tissues, and immune cells with attention to macrophages and monocytes?" <MRT-D-a>, "Produce all lysing enzymes and other cell chemistries required to destroy viruses and viral chemistries, break adverse host cell-virus linkages to cell structures and chemistries, and preserve the integrity of all cell cytoplasm and nucleus biochemical processes?" <MRT-D-c>,

j) "Any compromise of H/M-MB awareness of: Conditions of pathogen protein and RNA binding to cell chemical structures, microtubule transport systems, and the cellular rough endoplasmic reticulum and ribosome workbenches of chemical synthesis to co-opt cellular biochemical machinery for viral replication, support, and transport?" <MRT-S-a>, "Viral proteins or other chemistries that corrupt the body's clotting cascade resulting in extravascular hemorrhage and intravascular clotting?" <MRT-S-a>, "Glycoproteins or other pathogen chemistries that disrupt fluid systems transport and contribute to tissue edema, extravasation (vessel leakage) of blood and lymph, pneumonia, hypotensive shock, adverse CSF pressure?" <MRT-S-a>, "The capacity of cell biochemical systems to produce lysing enzymes and other chemistries to destroy

and/or silence viruses, virus-cell linkages, and viral counterfeit immune modulators and signaling chemistries?" <MRT-S-a>, "The capacity of cells to produce chemistries, energetic and vibrational cell actions to optimize cell signaling integrity, to prevent immune antibodies from being lost by their binding to cell-bound pathogen glycoproteins, to prevent viral attachment to membrane bound immune antibodies enabling cell infection?" <MRT-S-a>, "The capacity to energetically encourage antibody binding to vital areas of the pathogen for maximum lethality?" <MRT-S-a>, "Produce cell intelligence, energetic and chemical conditions to: efficiently block all disruptive effects of viral chemistries; to resist viral binding to cell structures, viral movement, and reproduction in the cell; to discriminate cellular from counterfeit pathogen immune modulators and signaling chemistries and respond only to authentic cell signals?" <MRT-D-c>,

k) "The presence on the bilipid viral envelope of all coronaviruses, a corona-like array of spike glycoproteins (S proteins) containing a receptor-binding domain S1 segment and a membrane-fusion S2 segment?" <MRT-S-a>, "The affinity of viral S proteins to bind to host cell membrane enzyme ACE2 as a receptor binding site, triggering the cell to encapsulate the virus by endocytosis and fuse the viral bilipid envelope and host cell membrane structures, introducing the virus to the cell?" <MRT-S-a>, "The behavior of the virus within the host cell – shedding its viral protein coat (capsid) and exposing its viral RNA structure, the transport of viral RNA to the host cell rough endoplasmic reticulum and Golgi apparatus where cell protein manufacturing resources are hijacked for viral replication?" <MRT-S-a>, "The release from the host cell by exocytosis of the newly replicated coronaviruses, and this cycle repeated with other host cells?" <MRT-S-a>, "Host cell capacity to recognize viral S protein structure and to produce host protective enzymes, antibodies, inert receptor binding domain agonists or other chemistries to damage the viral S proteins geometry and their receptor binding domain functionality?" <MRT-S-a>, "Optimize host cell intelligence to

recognize and compensate for these coronavirus mechanisms of cell infection?" <MRT-D-c>,

l) "Any compromise of awareness of the cellular apparatus for protein translation of any mRNA positioned upon cell ribosomes and rough endoplasmic reticulum, causing synthesis of the protein signified by that mRNA and its packaging for distribution by the Golgi complex?" <MRT-S-a>, "Potential protein synthesis from normal cell RNA, from the RNA of cancerous or otherwise diseased cells, and from viral RNA?" <MRT-S-a>, "The proteasome structure in the cytoplasm through which all translated proteins are passed and the resulting fragmentation of a small portion of each specific translated protein molecule into peptides characteristic of the protein from which they were cleaved?" <MRT-S-a>, "Host cell behavior to present each such peptide as an epitope attached to a cell membrane major histocompatibility complex protein (MHC) and constituting a record of all present-time host cell proteins production (human or viral)?" <MRT-S-a>, "T-cell monitoring and awareness of every body cell, and cell membrane presented host cell proteins and their significance representing normal cells, diseased cells, or viral infected cells, and T-cell coordination of an optimal immune response to viral infected, malignant, or severely diseased cells, and sparing of normal host cells?" <MRT-S-a>, "The capacity of the proteasome to fragment viral proteins into peptides representing proteins enfolded into the virus package of infected host cells and make them visible to T-cells and other immune cells?" <MRT-S-a>, "Optimize cellular mechanisms to present characteristic peptides to the adaptive immune system to direct an efficient response to viral infection, and to trigger apoptosis in cells whose peptide signatures indicate the cells is a threat to the body, and to direct protective behavior toward healthy host cells?" <MRT-D-c>,

m) "Any compromise of awareness of the cellular apparatus for protein translation of any mRNA that is positioned on to the ribosomes and rough endoplasmic reticulum for synthesis and packaged for distribution by the Golgi complex?" <MRT-S-a>, "Potential

protein synthesis from normal cell RNA, from the RNA of metaplastic or otherwise diseased cells, and from viral RNA?" <MRT-S-a>, "The proteasome structure in the cytoplasm through which all translated proteins are passed and the resulting fragmentation of a small portion of protein molecules into peptides characteristic of the protein they were cleaved from?" <MRT-S-a>, "The capacity of the proteasome to fragment viral proteins into peptides representing hidden viral proteins enfolded into the virus package and make them visible to T-cells and other immune cells?" <MRT-S-a>, "Cell behavior to attach each such peptide to a major histocompatibility complex protein (MHC) and install each to the plasma membrane to form an immune cell accessible record of all proteins (human or viral) that the cell is now making?" <MRT-S-a>, "T-cell monitoring of every body cell, searching for proteins recognized as normal cell, diseased or malignant cell, or infected cell viral proteins and to activate immune system mechanisms accordingly?" <MRT-S-a>, "Optimize cellular mechanisms to present characteristic peptides to the adaptive immune system for an efficient response to viral infection, and to trigger apoptosis in cells whose peptide signatures indicate the cells is a threat to the body?" <MRT-D-c>,

n) "The existence of coronavirus preferred infection routes to tissues and organs with high ACE2-expressing cells: type II alveolar cells (AT2) of the lung, upper esophageal stratified epithelial cells, absorptive enterocytes from ileum and colon, gall bladder and bile ductal cells, myocardial cells, kidney proximal tubule cells and bladder urothelial cells, the oral and nasopharynx mucosa and tongue epithelium?" <MRT-S-a>, "The capacity of cells to increase or decrease populations of cell membrane receptors and to produce inert agonists to these receptors to block coronavirus entry to the host cell?" <MRT-S-a>, "The presence of systemic and local tissue injury and pathophysiology from coronavirus infection and toxicity and the resultant insult to tissue hemodynamics, tissue fluid management, and inflammation with attention to: ARDS (94%), shock (44%), arrhythmia (38%), acute cardiac injury (26%), acute kidney injury (22%), and less commonly cerebral infarction, cerebral hemorrhage, and

hypoxic-ischemic encephalopathy?" <MRT-S-a> "Optimize host cell intelligence to limit coronavirus injury to host cells, tissues, and organs and to heal and restore the ideal physiological capacities to those tissues?" <MRT-D-c>,

o) **"Any introduction of bacteria or secondary pathogens in the wake of viral infection and disruption of immune defenses?" <MRT-D-a>, "Produce an efficient neuroimmune awareness, cell signaling, and response to eliminate all secondary pathogens?" <MRT-D-c>,**

p) **"The structures of self-awareness and physiological regulation of the nested hierarchy of the H-MB - CNS, ANSs including the lymphatic system of neurons and glia (astrocytes) and blood-brain barrier cells, and the M-MB - a nested hierarchy of many genera and many hundreds of microbial species and thousands of virus species - that engage in a dynamic real-time dialogue of awareness and immune, developmental, affective, mental, biorhythmic, and metabolic regulation?" <MRT-S-a>, The informational nested hierarchy of the H-MB - cell intelligence, CNS, ANSs, and of the M-MB - microbial cell intelligence, intraspecies and interspecies intelligence, and the overarching informational hierarchy of the H/M-MB super-organism?" <MRT-S-a>, "The production of H-MB and M-MB cell signaling via cell membrane release of signaling chemistries, and cellular resonant electromagnetic impulses and morphogenic field resonance among cells locally and at a distance?" <MRT-S-a>, "M-MB capacity to monitor and regulate populations of all organisms in the microbiota, with attention to the special role of the important spore-forming soil-based bacteria in this regulation?" <MRT-S-a>, "The H-MB and M-MB interpretation of metabolite levels in flora populated regions as chemical signaling of organ, system metabolic, and microbial conditions- particularly C2 acetic, C3 propionic, C4 butyric, and C5 valeric short chain fatty acids (SCFAs) metabolites?" <MRT-S-a>, "H-MB and M-MB monitoring of biological terrain conditions specific to local colonized tissues, with attention to pH, hydration, osmolality, oxygen gradient, and blood and lymph**

perfusion?" <MRT-S-a>, "The role of the microbiome to exchange with the neuroimmune system information regarding microbial conditions in the body for more efficient response to infection?" <MRT-S-a>, "Optimize H-MB and M-MB awareness of gut conditions and produce efficient manipulation of local conditions for greatest benefit to the H/M-MB super-organism?" <MRT-D-c>,

q) **"Cell intelligence awareness within the H-MB of their evolutionary history of interaction with viral and other microbial pathogens and flora?" <MRT-S-a>, "The capacity of human cell intelligence to recognize, distinguish, select, and to insert into the human cell genetic code the microbial genetic code sequences that are of potential value to enhance the human genome?" <MRT-S-a>, "The microbial origin of nearly half of human genetic code, 8% viral and 30% bacterial, and some level of other microbial sources. The capacity to recognize, distinguish, and resist the insertion of microbial genetic code sequences that are potentially harmful to the H-MB?" <MRT-S-a>, "The relatively higher level of microbial genetic code in metaplastic versus normal H-MB cells?" <MRT-S-a>, "Optimize H-MB cell awareness of the risk-reward potential to capture and insert specific microbial genetic code to exploit opportunities to enhance and broaden the H-MB cell genetic capacities, and to eliminate from the body all microbial organisms that endanger or otherwise burden H-MB cells?" <MRT-S-a>, "Optimize neuroimmune and cellular innate immunity physiology to accurately assess the risk to the H-MB of all viral and other microbial exposures and to efficiently modulate the immune response to avoid the self-injurious effect of positive feedback control loops, cytokine storm, and molecular mimicry, and to exploit any opportunities for H-MB advantage afforded by such exposures and interactions?" <MRT-D-c>,**

r) **"Any compromise of perfect awareness of the nested hierarchy of morphic fields comprising the growing field of human neuroimmune experience and wisdom accruing within the human population that has been infected by and successfully eliminated the current pandemic or epidemic infection, and avoided any self-injurious cytokine storm; or of the nested hierarchy of the current pandemic or epidemic**

infectious agents and the dendritic tree of mutagenic diversification into subspecies and transformation of their infectious properties?" <MRT-S-a>, "Optimize H/M-MB morphogenic attunement and awareness to the nested hierarchy of the current pandemic or epidemic and involved infectious agents and their subspecies and to the accruing morphic field of safe and successful human neuroimmune experience with these organisms to perfectly and safely direct a rapid and successful neuroimmune and innate cellular immunity systems response?" <MRT-D-c>, "Generate familiarity with the referenced infectious agents and the collective human neuroimmune response to them sufficient to produce a measured, proportional, timely, and efficient neuroimmune response, avoiding positive feedback neuroimmune dialog and resulting cytokine storm?" <MRT-D-c>,

s) MRT for the Y/N energetic representing the value in now applying CDRP, FNNP, ISOP, ARP in regard the referenced viral and other pathogens. Use the EVCP Worksheet.

3) **"Shall the H/M-MB now: Project its awareness, attention, belief, and intention into all times and dimensions of the undivided universal field of consciousness and identify, accept, and release attunement to all events, IFs, and IFs-h (IFs that hide subject IFs from awareness) that define into existence the transformed-from system state?" <MRT-S-a>, "Or, that cause this transformation to seem dangerous, undeserved, or impossible to achieve? <MRT-S-a>, "Create awareness, acceptance, attunement, and belief of all corrective events and TrIQs to define into existence the transformed-to system state?" <MRT-S-a>, "Create the intention to collapse the quantum potential of the field's reality quanta to manifest this transformation?" <MRT-S-a>, "Create optimal attunement and coherence to the source of creation, innate intelligence, and the consciousness of healing and accept the role of co-creator of all experienced reality, NOW?" <MRT-D-c>.** Refresh this awareness and intention with each MRT pulse. The dynamic MRT end point signals that the survey is complete, awareness is optimal, and this transformation is now informationally realized.

4) Optionally: **“After fully applying the therapeutic intention of this correction, H/M-MB projection of the TSB metric for the TFS is $\log_{10}$1000 L- ___?” <MRT-S-a>,**

   a) If the resulting projected TSB is not acceptable: **“Would it be of benefit to now reapply the therapeutic intention of this correction 1000 times in the next 10 seconds?” <MRT-D-c>.** If yes, make a correction, holding this intention and reapplying the UCS.

   b) Recheck the question, **“After fully applying the therapeutic intention of this correction, H/M-MB projection of the TSB metric for the TFS is $\log_{10}$1000 L- ___?” <MRT-S-a>. If MRT suggests it would be of value, make the correction again. “Would it be of benefit to now reapply the therapeutic intention of this correction 1000 times in the next 10 seconds?” <MRT-D-c>**, and so on, until an acceptable level of correction has been reached.