

# Chimpanzees, Genes, and



## EPIGENETICS:

If we encounter evidence from science that challenges the biblical creation account, how should we respond? As Adventist educators, we can't shield our students from challenging information, but we have a responsibility to guide them to think clearly for themselves and to critically evaluate what they read or hear.

Some scholars allow evolutionary science to guide their understanding of theology, attempting to make their theology fit the latest scientific paradigms. But recent episodes in science remind us that this is not the wisest approach. Even scientific research publications are often challenged within a short time by the findings of other researchers, and this can be seen in the subject of evolution. Science is a fascinating and productive process of discovery, but it is a very human endeavor and has its limits, especially in the scientific study of ancient history.<sup>1</sup>

In the mid-1800s, a living cell was thought to be like a bit of jelly with a few things floating in it. Scientists viewed life as based on a simple cell that could change and evolve easily along the path toward complex life. Since then, we can point to well-known events that mark significant growth in under-

## Changing Views of Inheritance

standing of life. Gregor Mendel's experiments with garden peas (1856-1863) laid a foundation for the field of genetics, although it took several decades to become well known. This was followed by the neo-Darwinian synthesis (or modern synthesis, basically the union of Mendelian genetics, Darwinian natural selection, and paleontology), which remains the predominant

theoretical influence for evolutionary biologists.<sup>2</sup>

Another milestone was James Watson and Francis Crick's 1950s discovery of the structure of the DNA molecule. After that, it seemed that human beings now understood the instruction book that determined all the structures of cells and living organisms, and what directed the process of development from an egg to a mature animal. But that was a premature conclusion.

### Advances in Biochemistry and Molecular Biology

We could refer to the past half century or so as the era of biochemistry and molecular biology, with its exponential growth in knowledge about the basis of life. Over the past 30 years, scientists have discovered that the complexities required for carrying out the necessities and vibrant activities in each cell are astound-

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ing. Much of what is happening inside of a cell can't be directly observed but must be ferreted out with clever biochemical detective work. It has been enjoyable to attend research seminar presentations by molecular biologists and to learn about the increasingly insightful logic and laboratory procedures they use to discover what is going on in the recesses of a minute living cell.

This detective work by the scientific community has periodically received significant boosts from new analytical equipment such as high-speed ultracentrifuges and spectrophotometers, and combined with new techniques that fuel advances in understanding, enable scientists to dig deeper into the complex structures and processes of life. The use of gene-sequencing procedures has enabled scientists to discover the sequence of nucleotides in DNA in order to define the exact information in the genetic library of each organism. The mapping of the complete sequence of the human genome by 2004 was an impressive breakthrough.<sup>3</sup>

In time, it became evident that having the human genome sequence was only the tip of the iceberg. In fact, in some ways, we are just beginning to understand the biochemical symphony that is constantly underway in every living cell. Until fairly recently, it was believed that the instructions in DNA provided one-way control over the processes in the cell. DNA directed the formation of RNA, which made proteins. This concept was considered the central dogma of molecular biology.<sup>4</sup> Scientists identified two types of genes, *coding genes* with the instructions (or code) for making specific proteins, and *non-coding genes*, which were sometimes called "silent genes" because they didn't seem to do anything. Many of the non-coding genes were considered "junk DNA"—functionless leftovers from the process of evolution that could mutate and evolve into new genes, producing new proteins and eventually new structures and physiological processes in the course of evolution.

Then researchers discovered another category of genes—regulatory genes, which regulate or control the coding genes by encoding a protein, RNA, or microRNAs (mRNAs). There is much more to that control process, but let's not get ahead of our story.

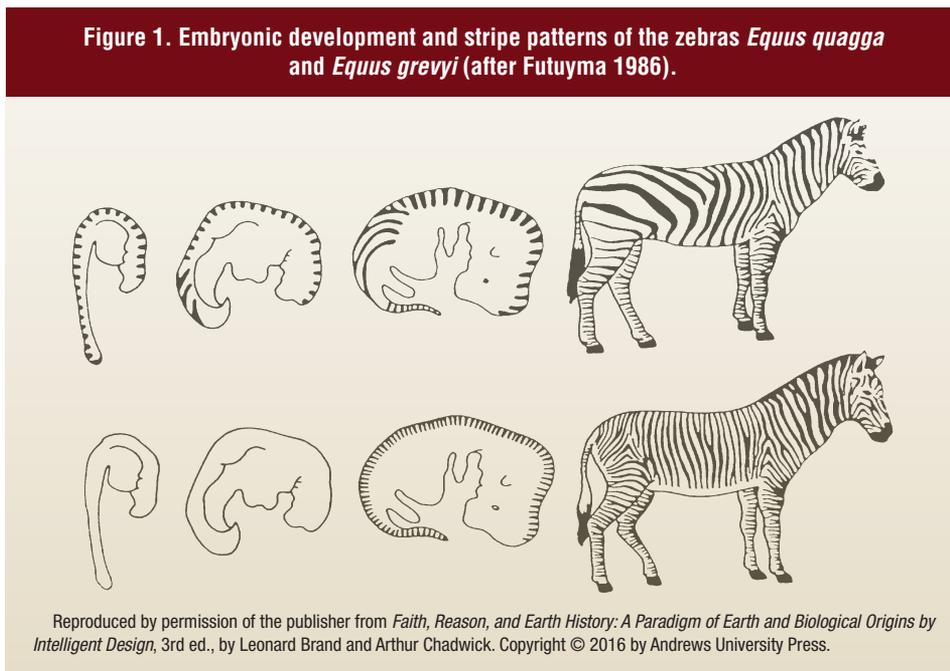
### Regulatory Genes and "Junk DNA"

Small influences on the actions of regulatory genes can result in large changes in an animal, including controlling the timing of embryological development and thereby changing the phenotype. A simple example appears in the study of zebras. Timing of embryonic events apparently controls the stripe pattern in some zebras. The black-and-white stripe patterns on the lower back of the zebra *Equus quagga* are stretched out and

unevenly spaced. This results because the back part of the embryo grows faster than the rest of it after the pattern is established. The stripe pattern in Grevy's zebra, *Equus grevyi*, is not established until after that differential growth is completed. Consequently, the stripes in the adult of this species are more evenly spaced (see Figure 1).<sup>5</sup> This change in the developmental process is known as heterochrony, which is simply a change in the timing of a developmental process.

Other discoveries have contributed to improved understanding. The human genome is 98 percent non-coding DNA. If this is all junk DNA, it would imply that very little of our genome is functional. But over the years, regions of DNA formerly thought to contain junk DNA have been found to be regulatory regions, which control the action of coding genes.<sup>6</sup> This change in thinking reached a turning point in September

**Figure 1. Embryonic development and stripe patterns of the zebras *Equus quagga* and *Equus grevyi* (after Futuyama 1986).**



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of 2012 at the completion of the massive government-funded ENCODE genetic-research project. After intensive study of the human genome, some 30 papers were published, which recognized that most of the formerly tagged "junk DNA" is functional and consists of types of regulatory regions.<sup>7</sup> There are still vast unknowns regarding the genetic system, and future research may show that essentially all DNA is functional.

Some evolutionary scientists have criticized the ENCODE conclusions,<sup>8</sup> but their objections are based on evolutionary theory, not on observation. In deciding how much credence to give such criticisms and their underlying theory, remember that science is always a continuing search for understanding, and it is often wise to wait for new data to provide better answers to our questions, rather than concluding that a particular discovery is the "final word."

All those regulatory genes and the biochemical pathways associated with them are not only functional, but important in controlling how, when, and where the coding genes make their

proteins and how each protein integrates with other “building materials” in the cell. This can be compared to how engineers determine when and where to use bricks and other building material in a construction project. The same bricks can become part of a doghouse or a palace. The engineers are the regulatory system that decides how to use bricks to build the appropriate structure. In living cells, the regulatory genes, formerly interpreted as junk DNA, are the engineers that manage the activity of coding genes. This is truly a paradigm shift in progress. As a result, “junk DNA” is no longer a useful concept.

### Epigenetics and DNA Functioning

In recent years, rapidly advancing knowledge in biochemistry and molecular biology has been forcing a re-evaluation of evolutionary theory. Information-bearing molecules like hormones, DNA, and proteins are astonishingly complex. Rather than a single (DNA) information system, the genetic system has in recent years become vastly more complex because of the discovery of cell processes referred to as *epigenetics*. The blossoming field of epigenetics is revolutionizing genetics.<sup>9</sup> Epigenetics, which means “above the genes,” is the study of processes outside of DNA that affect how DNA information is interpreted.

DNA code contains vast amounts of information, but layers of additional control systems manage and regulate how the DNA is expressed. DNA is like a hard drive containing a massive amount of information, but this information is managed by the epigenetic system. DNA cannot do anything by itself. It requires the epigenetic complex of RNAs, proteins, etc. to direct and control how the DNA information is to be interpreted as it is put to work in the cell.<sup>10</sup>

The epigenetic system does not just read and blindly apply the DNA information. It uses information that the organism’s sensors have detected from its environment, and combines this with the DNA information to determine how to use the DNA to respond appropriately to the environmental cues.<sup>11</sup> In this process, millions of chemical side groups in each cell, including methyl and acetyl, can be attached to specific places on DNA and associated molecules, and serve as chemical tags or markers that turn genes on or off, controlling whether each gene will be active, and if so, when and how. Genes are not always completely on or off—the tags also act like dimmer switches, varying the genes’ level of activity. Additional levels of intracellular management control where and when these tags will be associated with the DNA.

A great example of epigenetic control involves the hormone cortisol. The reduction in this stressor in domesticated dogs has a large impact on the difference between domesticated dogs and wild wolves or foxes because of its involvement in these animals’ development. At a Silver Fox farm in Soviet Russia, Dmitry

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Belyaev, a Russian scientist, wanted to develop a more easily managed breed of foxes.<sup>12</sup> Belyaev allowed only certain foxes to breed, selecting for only one factor, signs of tameness, i.e., lack of fear or aggressiveness toward humans. After only 10 to 18 generations, the foxes had changed their behavior and were much tamer. But this was only the beginning! They had also changed physically and physiologically. They developed floppy ears, shortened, curled tails, as well as a more dog-like snout (broader and shortened), piebald coat color (a black-and-

white pattern), and began to sound like dogs. Some females came into estrus twice a year instead of once, as in wild foxes.

Why these changes? Since the *only* thing that was originally sought after was tameness, why the other domestic dog-like changes? Belyaev’s selection for tameness changed the level of cortisol, a stress hormone. He found that the formation of cortisol was delayed during development in the “tamer” foxes, which played an important role in development. This resulted in the retention of certain fox “puppy” features in the adults (paedomorphosis), including floppy ears and curled tails. The puppies were also much more likely to seek out human companionship, due to a youthful lack of adult stress responses. The animals also became sexually mature at a younger age. All of this equals heterochrony—a difference in the timing of developmental processes. These changes did not result from mutations but from a change in how the DNA information was interpreted and applied. We are all familiar with the extreme variation in body shapes and other features in breeds of dogs, which is interpreted as the result of a similar process to that seen in the fox experiment.<sup>13</sup>

The complexity and details of this system are just beginning to be understood. Epigenetic processes have the amazing quality of allowing environmental cues to influence changes in cells and organisms without any change in their DNA.<sup>14</sup> Epigenetic tags alter the activity of certain genes, and these alterations have a distinct influence on physiology, anatomy, and even behavior of animals. The DNA doesn’t change, but epigenetic tags control how the information in DNA is used or not used. The old idea of genetic instructions going only from DNA to RNA to protein, but never in the other direction, is now known to be false. As one molecular biologist has described it, the amazing system inside of a cell “decides” how the DNA will be expressed.<sup>15</sup>

### Unto the Third and Fourth Generation

These intracellular decisions, influenced by environmental cues, can be passed on to offspring for several generations. We can propose that this incredible epigenetic system was created to allow living organisms to adjust to environmental condi-

tions. However, when the environment contains destructive elements, as it often has since the entrance of sin into our world, the epigenetic results can be unpleasant.

Stress, nutrition, and other aspects of a mother's experiences during pregnancy can alter her young through epigenetic processes, with the results lasting through the offspring's lifetime, and can even affect several succeeding generations.<sup>16</sup> This brings to mind the Bible statement "punishing the children for the sin of the parents to the third and fourth generation."<sup>17</sup> A recent epigenetics article in the prestigious scientific journal *Nature* was even entitled "Epigenetics: The Sins of the Father."<sup>18</sup>

An example of how epigenetics affects offspring for several generations comes from World War II. The Dutch Hunger Winter, or the Dutch Famine, occurred near the end of the war in 1944-1945, when German military took revenge on the uncooperative Dutch people by blockading part of their country during most of that winter, preventing them from receiving food or other supplies. More than 20,000 people starved to death, and the rest of the population was reduced to a starvation diet containing as little as 30 percent of the nutrition needed by a human body.

National health records revealed that children born during or soon after the Dutch Famine had specific health defects, depending on what portion of their embryonic development occurred during the famine.<sup>19</sup> If a mother experienced the famine during the last few months of pregnancy, her baby was born small, remained small all of his or her life, and had a lower chance of obesity than normal. These children never fully recovered from their poor pre-birth nutrition. Those mothers who experienced the famine only during the early months of pregnancy usually had normal birth-weight babies who experienced higher-than-normal rates of obesity later in life as well as other health problems. Some of these effects were evident even into the second generation. Although the grandchildren of the malnourished mothers<sup>20</sup> did not experience the famine, they were affected by the epigenetic carryover from a previous generation.

An experiment with mice revealed epigenetically inherited behavior persisting for more than one generation. Mice were exposed to the mild odor of acetophenone, a chemical with a sweet smell. At each exposure to this odor, they received a mild shock to their foot several times a day for three days. They became fearful and would freeze whenever they smelled the odor, even if they did not receive a shock.

The mice in the experiment were then mated with females that had never been exposed to acetophenone. Their offspring, when mature, were exposed to the same

odor but without any shock. The young mice were more sensitive to this odor than other odors, and were generally startled by any noise when exposed to the odor. Their offspring, the grandchildren of the initial experimental mice, showed the same behaviors. All three generations had an enlargement of the brain structure where the odor-sensitive neurons connect with the olfactory bulb. This supports the conclusion that epigenetic inheritance can persist through several generations.<sup>21</sup>

In another well-documented study, exposing mice to an environmental toxin produced effects on epigenetic inheritance for several generations. Pregnant laboratory mice were given a dose of the fungicide vinclozolin. Their male offspring were born with testicular defects and reduced fertility. These effects lasted for at least four generations, despite no further exposure of the mice to the toxin. Research showed that the defects did not result from DNA mutations but were epigenetic effects.<sup>22</sup>

In some cases, epigenetics has been shown to underlie microevolutionary changes, including cyclic variation in the size of beaks of Galapagos finches in response to variations in the available food supply.<sup>23</sup> Also, blind fish living in dark caves are not sightless because of mutations to their eye genes, which are all functional. The blindness is caused by epigenetic processes that turned off the eye genes.<sup>24</sup>

### A Challenge to Neo-Darwinism

Evolutionary scientists have been reticent to accept the notion that environmental factors could influence inheritance because it is difficult to reconcile this with Neo-Darwinian theory. For inheritance to be affected by the environment seemed like Lamarckism, in which an organism passes on characteristics acquired during its lifetime (an idea proposed by Jean Baptiste Lamarck, a contemporary of Charles Darwin). This concept is not compatible with Neo-Darwinism, but accumulating evidence supports the reality of heritable environmental effects on the genome.

Why is this a problem for evolution to explain? If life forms arose with no designer or Creator, all new biological changes must begin as the result of random mutations, followed by natural selection. The mutation process must be random, meaning that it cannot comprehend what the animal needs, or it would be apparent that someone, somewhere in the process, knows what changes need to occur. But the new findings indicate that beneficial, inheritable changes can be initiated by environmental influence (now called Neo-Lamarckism), further indicating that the genetic/epigenetic system is designed to recognize, in a non-random way, what

### Recommendations for Further Reading

The following sources are recommended for deeper study of the issues discussed in this article:

Leonard Brand and Arthur Chadwick, *Faith, Reason, and Earth History: A Paradigm of Earth and Biological Origins by Intelligent Design*, 3rd ed. (Berrien Springs, Mich.: Andrews University Press, 2016), specifically chapters 8 and 10.

Leonard R. Brand and Richard M. Davidson, *Choose You This Day: Why It Matters What You Believe About Creation* (Nampa, Idaho: Pacific Press Publ. Assn., 2013).

Nessa Carey, *The Epigenetics Revolution* (New York: Columbia University Press, 2012).

Richard C. Francis, *Epigenetics: The Ultimate Mystery of Inheritance* (New York: W. W. Norton and Co., 2011).

Stephen C. Meyer, *Darwin's Doubt: The Explosive Origin of Animal Life and the Case for Intelligent Design* (New York: HarperOne, 2013).

changes will be beneficial. Evolutionary scientists generally ignore this evidence,<sup>25</sup> or discount it, or assume that somehow they will be able to explain it.

### Natural Genetic Engineering

James Shapiro, an eminent molecular biologist and committed evolutionary scientist, recently published a book entitled *Evolution: A View From the 21st Century*.<sup>26</sup> His message is that molecular biology research in the past few decades has shown a level of complexity and cellular sophistication that is incompatible with Neo-Darwinian random mutations and natural selection as a means of change in living organisms, since it extends beyond minor variations. When sensors in the cell detect environmental cues, the molecular complex within the cell uses these cues to make decisions about how to interpret the DNA information and respond to the environment. Shapiro calls this natural genetic engineering and regards the evolution of intracellular system as a mystery, along with the origin of life. Traditional Neo-Darwinians criticize him, but he says, “their criticisms are philosophical, not scientific,”<sup>27</sup> and lack empirical evidence and support. Their reliance on random mutations as the engine for evolution does not stand up to the growing evidence. The cell, Shapiro concludes, is too sophisticated for random mutations to be an effective process for evolutionary change. He has not become a creationist, but he is honestly trying to deal with the evidence in ways that are compatible with his worldview.

### What Chimpanzee Genes Can Tell Us About Theology

This growth of knowledge about genetics brought with it an episode that can help us understand the proper relationship between science and the Bible. In 2005, after scientists sequenced the chimpanzee genome, some geneticists determined that chimpanzee genes and human genes were 98 to 99 percent similar (comparing only genes that are found in both species).<sup>28</sup> It would seem, then, that humans are hardly any different genetically from a chimpanzee, and evolutionists asserted that this proved that humans and chimps had obviously descended from common ancestors.

How could Christians argue against this evidence? Even before the final analysis of the chimpanzee genome, books were being published that struggled with the implications for Christians of this 98-percent similarity.<sup>29</sup> Many Christians accepted the discovery as further evidence against biblical creationism, but it turned out to be an instructive case study in how to respond to troublesome evidence.

Sometimes it is wise to deal with such evidence by just waiting for future discoveries because the ongoing progress of science very often has surprises for us. That was a wise approach in this case. The first clue came from considering the context of that 98-percent similarity. The original study of the chimpanzee genome in 2005 was not based on the entire genome, but only a selected portion, and was done in a way that emphasized genes that code for protein. Thus, researchers studied only a fraction of the genome, the part that

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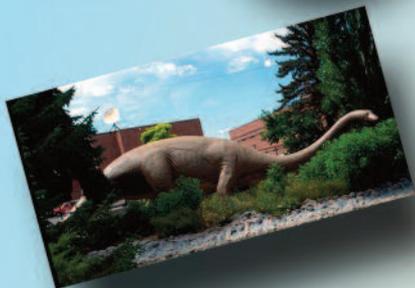
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is more similar. However, the part of the genome that is expected to show the important differences would be the former junk DNA, which includes the regulatory part of the DNA.<sup>30</sup> Also, a later comparison of the human and chimpanzee male Y chromosome (the male sex chromosome) revealed that this chromosome in the two species “differ[s] radically in sequence structure and gene content.”<sup>31</sup>

As we discussed above, scientists found that all this non-coding DNA was not junk at all, but important, functional, regulatory, or structural nucleotides.<sup>32</sup> Rather than the human genome being mostly functionless junk, it is actually a massive, sophisticated control system with a complex of proteins and various types of RNA and chemical tags that direct the activity of the protein-coding genes. These epigenetic factors tell when, where, and how much of each protein is to be produced, and how they will connect to one another to make the parts of cells and of animal bodies. How naïve of us not to have realized long ago that such a control system was essential.

Similar bricks can be used to make a doghouse or a palace. In a similar way, proteins can be used to make a human, a chimpanzee, or a mouse depending especially on the instructions provided by the regulatory genes and epigenetics. A striking example of this principle has been demonstrated with the *Pax6* gene that manages the development of the eye. Genes like this, which control development, are common to all Animalia. When the *Pax6* gene is removed from a mouse and placed in the area of fly DNA where the jaw of *Drosophila* (a fruit fly) is supposed to form, it causes the formation of a compound fly-eye—not a jaw or a mouse eye but a fly-eye!<sup>33</sup> It is obvious that this gene is so similar in mice and flies that the developmental system in the fly recognizes the mouse gene as if it were a fly gene. But since the genetic system in the fly is designed to make fly parts, not mouse parts, the fly cells make the eye of a fly.

In like manner, the human genetic/epigenetic system is designed to use its proteins to make a human, not some other animal. We could expect it to still make a human, even if our protein coding genes were 100 percent similar to chimp genes, which they aren't. These new genetic insights help explain how humans can be radically different from chimpanzees in spiritual, mental, and fine motor skills in spite of the similarity of our protein-coding genes.

There is a significant lesson in this, for our view of theology. In 2005, we could have decided that our theology must accommodate the genetic evidence for human and chimpanzee evolution from common ancestors, as many Christians have done. Or we could base our theology on the Bible, stay with the Genesis creation account, and predict that new scientific discoveries will resolve more of our uncertainties.

If we take the first approach and make contemporary scientific interpretations our standard, science is likely to move on and leave our theology without a foundation, just like the demise of junk DNA, which, within a period of a decade, left an empty place in the theories of those who took current discoveries too seriously. Additional scientific findings are building a stronger case for accepting creation.

Conversely, those who were willing to wait patiently for new insights in genetics were rewarded by discoveries that greatly reduced the proposed reasons for accepting evolutionary conclusions about human and chimp relationships.

Today, other aspects of science are used to support the theory of all of life evolving from common ancestors, but we suggest that it is wise to wait to see what science will discover next, since the biological evidence for evolution of new organisms or physiological systems is running into increasing difficulty.<sup>34</sup>

Darwin's theory was developed at a time of great biological naïveté combined with a rebellion against religious authority.<sup>35</sup> Parts of the theory of evolution, including microevolution and speciation, are still valid and well supported. But modern discoveries in genetics and epigenetics are introducing more and more serious problems for the theory that major groups of organisms have arisen by evolution. However, evolution textbooks and technical scientific publications continue their persistent, authoritarian claims that life arose without a designer and diversified by processes of naturalistic evolution.<sup>36</sup> It is becoming increasingly evident that these claims depend more on evolutionary or naturalistic assumptions than on evidence. More than ever before, it is not sufficient to just read and accept published claims, especially in assumption-dominated subjects like origins. It is imperative that we encourage our students to learn to critically evaluate scientific interpretations before accepting them. ✍

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- Leonard Brand, "Naturalism: Its Role in Science," op. cit.
- Brand and Chadwick, *Faith, Reason, and Earth History*, op. cit.

Teachers are to “. . . consider the highest good of their students as individuals, the duties that life will lay on them, the service it requires, and the preparation demanded.”<sup>8</sup> This is not a humanly directed task; it requires partnership with the divine—God, through the Holy Spirit. To partner with the divine is a “noble work”<sup>9</sup>—one of building character, both teacher and student, and it is work that “cannot be completed in this life, but will be continued in the life to come.”<sup>10</sup>

And yet, this is what we as Seventh-day Adventist Christian educators are called to do every day! Individually and collectively, we engage in thinking about how best to prepare our students to live in this world, while anticipating a future world. When we look at our students, do we see them as they are, or as what they can become? Do we prepare them for the world in which they live, or the world they will inhabit 10, 15, or 20 years into the future? What skills and dispositions will we need to focus on so that we can empower them to succeed? And, when we look in the mirror, what will we do with what we see reflected there?

The task may seem difficult and overwhelming; however, we have help. We have a cheerleader who says “You can do it!” In Jeremiah, we’re told: “Call to Me, and I will answer you” (Jeremiah 33:3 NIV),<sup>11</sup> and in James we’re reminded that “If any of you lacks wisdom, you should ask God, who gives generously to all” (James 1:5, NIV). From navigating the challenges of teaching in a postmodern world (John Wesley Taylor V) to following after the Master Teacher (Laszlo Gallusz) or wrestling with scientific discovery (Leonard Brand and Carl Person), we have the assurance of divine assistance. And when implementing new strategies or programs (Barbara Fisher, R. Lee Davidson and Tammy Overstreet, and Renard and Jayne Ann Doneskey), we have the assurance of wisdom and insight. Above all, when interacting with our students, let’s see them as God sees us now and as Mr. Gatewood saw the countless children that crossed his path each day—as characters fit for the kingdom. With

whole-souled, large-hearted devotion, we can each continue to grow in Him, and “. . . more fully to reflect . . . the light of the knowledge of His glory”<sup>12</sup> for eternity.

#### NOTES AND REFERENCES

1. Guy Larry Gatewood taught 5th and 6th grade at Ruth Murdoch Elementary School, Berrien Springs, Michigan.
2. Ellen G. White, *True Education* (Nampa, Idaho: Pacific Press Publ. Assn., 2000), p. 171.
3. The significance of not only remembering names, but also pronouncing them correctly, is not lost on students and contributes to their sense of belonging and acceptance within the community of learners. The act of taking the time is what counts. See Corey Mitchell’s commentary, “A Teacher Mispronouncing a Student’s Name Can Have a Lasting Impact,” *The Rundown* (May 16, 2016): <http://www.pbs.org/newshour/rundown/a-teacher-mispronouncing-a-students-name-can-have-a-lasting-impact/#.Vzs1c5IEbhE.mailto>. Another useful resource is <https://www.name-coach.com/>, a program designed to assist in name pronunciation.
4. Volunteering in Mr. Gatewood’s classroom helped me complete the Pathfinder Investiture Achievement program: <http://www.pathfinderonline.org/classes/overview>. I later worked as his reader until graduating from high school.
5. Parker J. Palmer, *The Courage to Teach: Exploring the Inner Landscape of a Teacher’s Life* (San Francisco: Jossey-Bass, 2007), p. 2.
6. *Ibid.*, p. 3.
7. White, *True Education*, op. cit., p. 174.
8. *Ibid.*, p. 13.
9. *Ibid.*
10. Ellen White’s expansive thoughts on the value of seeing our students as what they might become are well stated in *Counsels to Parents, Teachers, and Students* where she encourages every teacher to “see in every pupil the handiwork of God—a candidate for immortal honors. He [the teacher] should seek so to educate, train, and discipline the youth that each may reach the high standard of excellence to which God calls him [or her]” (Mountain View, Calif.: Pacific Press Publ. Assn., 1943), p. 229.
11. Jeremiah 33:3; James 1:5. All Scripture quotations are taken from the Holy Bible, New International Version®, NIV® Copyright © 1973, 1978, 1984, 2011 by Biblica, Inc.® Used by permission. All rights reserved worldwide.
12. Ellen G. White, *Education* (Mountain View, Calif.: Pacific Press Publ. Assn., 1903), p. 22.

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